

A Beta Way: A Tutorial For Using Beta Regression in Psychological Research

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24

Abstract

25 Rates, percentages, and proportions are common outcomes in psychology and the social sciences.
26 These outcomes are often analyzed using models that assume normality, but this practice
27 overlooks important features of the data, such as their natural bounds at 0 and 1. As a result,
28 estimates can become distorted. In contrast, treating such outcomes as beta-distributed respects
29 these limits and can yield more accurate estimates. Despite these advantages, the use of beta
30 models in applied research remains limited. Our goal is to provide researchers with practical
31 guidance for adopting beta regression models, illustrated with an example drawn from the
32 psychological literature. We begin by introducing the beta distribution and beta regression,
33 emphasizing key components and assumptions. Next, using data from a learning and memory
34 study, we demonstrate how to fit a beta regression model in R with the Bayesian package {brms}
35 and how to interpret results on the response scale. We also discuss model extensions, including
36 zero-inflated, zero- and one-inflated, and ordered beta models. Basic familiarity with regression
37 modeling and R is assumed. To promote wider adoption of these methods, we provide detailed
38 code and materials at <https://doi.org/10.5281/zenodo.15830595>.

39 *Keywords:* beta regression, beta distribution, R tutorial, psychology, learning and memory

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Introduction

Many outcomes in psychological research are naturally expressed as proportions or percentages. These include the proportion of correct responses on a test (e.g., Kornell and Bjork (2008)), the proportion of time a participant fixates on a stimulus in an eye-tracking task (e.g., James et al., 2025), or the proportion of respondents endorsing a particular belief (e.g., Costello et al., 2024). A central question for studies involving proportional data is how to analyze these outcomes in a statistically appropriate and interpretable way.

Researchers frequently default to linear models that assume Gaussian (normal) distributions, such as t-tests, ANOVAs, and linear regression. However, these models make strong assumptions: (1) residuals are normally distributed, (2) the outcome is unbounded (from $-\infty$ to ∞), and (3) variance is constant across the range of the data. These assumptions are rarely satisfied in practice (Sladekova & Field, 2024), and they are especially ill-suited for proportional outcomes, which are bounded between 0 and 1 and often exhibit heteroscedasticity—non-constant variance, particularly near the boundaries (Ferrari & Cribari-Neto, 2004; Paolino, 2001; Smithson & Verkuilen, 2006). Violating these assumptions can lead to biased estimates and spurious inferences, especially when many observations cluster near 0 or 1.

In some cases, a generalized linear model (GLM) can relax the assumption of normality. For example, binomial and Bernoulli models—often referred to as logistic regression when used with a logit link—are appropriate when the outcome is binary (0/1) or represents the number of successes out of a fixed number of trials. These models, however, rely on data that arise from discrete trial structures. They tend to perform poorly when the observed proportions are truly continuous or when the data show extra variability (overdispersion), particularly when many

24 values occur near the boundaries of the scale (0 and 1).

25 The challenges of analyzing proportional data are not new (see Bartlett, 1936).

26 Fortunately, several existing approaches address the limitations of commonly used models. One
27 such approach is beta regression, an extension of the generalized linear model that employs the
28 beta distribution (Ferrari & Cribari-Neto, 2004; Paolino, 2001). Beta regression offers a flexible
29 and robust solution for modeling proportional data directly by accounting for boundary effects and
30 over-dispersion, making it a valuable alternative to traditional binomial models. This approach is
31 particularly well-suited for psychological research because it can handle both the bounded nature
32 of proportional data and the non-constant variance often encountered in these datasets (Sladekova
33 & Field, 2024). In addition, the direct modeling of proportions allows comparability across tasks
34 and scales, and can be particularly valuable when only the proportional data is available, as is
35 often the case with secondary data that lack item-level structure or point values. While usage of
36 these models has faced obstacles due to theoretical and computational limitations, as we argue in
37 this paper, beta regression and its extensions now provide an accessible and more robust method
38 to traditional linear modeling.

39 While in this paper we will focus on proportional-responses that lie between 0 and 1—it is
40 important to note that our analysis applies to any bounded continuous scale. Any bounded scale
41 can be mapped to lie within 0 and 1 without resulting in a loss of information as the
42 transformation is linear.¹ Consequently, a scale that has natural end points of -1,234 and
43 +8,451—or any other end points on the real number line short of infinity—can be modeled using
44 the approaches we describe in this paper.

45 A Beta Way Is Possible

46 With the widespread availability of open-source software such as R (R Core Team, 2024)
47 and its extensive ecosystem of user-developed packages, advanced models like beta regression
48 have become increasingly accessible to applied researchers. Yet, their adoption in psychology

¹ Specifically, for any continuous bounded variable x , we can rescale this variable to lie within 0 and 1 by using the formula $x' = \frac{x - \min(x)}{\max(x) - \min(x)}$ where $0 \leq x' \leq 1$.

49 remains relatively limited. One contributing factor may be the lack of domain-specific examples
50 that demonstrate how these models address common challenges in psychological data. Although
51 recent years have seen a growing interest in beta regression, and a number of useful tutorials are
52 available (Bendixen & Purzycki, 2023; Coretta & Bürkner, 2025; Heiss, 2021; e.g., Smithson &
53 Verkuilen, 2006; Vuorre, 2019), these resources often either focus on basic implementation or
54 briefly mention extensions without detailing how they can be applied to psychologically relevant
55 research questions.

56 The present tutorial aims to help bridge this gap by providing a comprehensive, applied
57 introduction to beta regression and several of its extensions. In addition to the standard beta
58 model, we walk through zero-inflated, zero-one-inflated, and ordered beta regression. These
59 models are particularly useful for researchers working with proportion outcomes that include
60 boundary values (e.g., exact 0s or 1s) or responses with an inherent ordinal structure. Our goal is
61 to offer practical guidance that enables psychological researchers to implement, interpret, and
62 report these models in ways that directly support their empirical questions.

63 Beyond model specification, we place strong emphasis on interpreting results on the
64 response scale—that is, in terms of probabilities and proportions—rather than relying on often
65 difficult to interpret parameters. This focus makes the models more accessible and meaningful for
66 psychological applications, where effects are often easier to communicate when framed on the
67 original scale of the outcome (e.g., changes in recall accuracy or task performance). Throughout,
68 we provide reproducible code and annotated examples to help readers implement and interpret
69 these models in their own work.

70 We begin the tutorial with a non-technical overview of the beta distribution and its core
71 parameters. We then walk through the process of estimating beta regression models using the R
72 package `{brms}` (Bürkner, 2017), illustrating each step with applied examples. To guide
73 interpretation, we emphasize coefficients, predicted probabilities, and marginal effects calculated
74 using the `{marginaleffects}` package (Arel-Bundock et al., 2024). We also introduce several useful
75 extensions—zero-inflated (ZIB), zero-one-inflated (ZOIB), and ordered beta regression—that enable

76 researchers to model outcomes that include boundary values. Finally, all code and materials used
77 in this tutorial are fully reproducible and available via our GitHub repository:
78 https://github.com/jgeller112/beta_regression_tutorial and on Zenodo
79 (<https://doi.org/10.5281/zenodo.15830595>)².

80 **Beta Distribution**

81 Proportional data pose some challenges for standard modeling approaches: The data are
82 bounded between 0 and 1 and often exhibit non-constant variance (heteroscedasticity) (Ferrari &
83 Cribari-Neto, 2004; Paolino, 2001). Common distributions used within the generalized linear
84 model frameworks often fail to capture these properties adequately, which can necessitate
85 alternative modeling strategies.

86 While we do not have time to delve fully into its derivation, the beta distribution is useful
87 for modeling bounded continuous scales because it is the distribution for the probability of an
88 event. Given that a probability can take on any value from near 0 (the event will not occur with
89 certainty) to 1 (the event will occur with certainty), the beta distribution can likewise take on
90 virtually any value in that bounded interval. As a consequence, the beta distribution is the
91 maximum entropy distribution for *any* bounded continuous random variable, which means that the
92 beta distribution can represent the full range of possibilities of such a scale.³ As a consequence, if

² In this article, we try to limit code where possible; however, the online version has all the code needed to reproduce all analyses herein. Furthermore, to promote transparency and reproducibility, the tutorial was written in R version 4.5.1 (R Core Team (2024)) using Quarto (v.1.5.54), an open-source publishing system that allows for dynamic and static documents. This allows figures, tables, and text to be programmatically included directly in the manuscript, ensuring that all results are seamlessly integrated into the document. In addition, we use the `rix` (Rodrigues & Baumann, 2025) R package which harnesses the power of the `nix` (Dolstra & contributors, 2006) ecosystem to help with computational reproducibility. Not only does this give us a snapshot of the packages used to create the current manuscript, but it also takes a snapshot of system dependencies used at run-time. This way reproducers can easily re-use the exact same environment by installing the `nix` package manager and using the included `default.nix` file to set up the right environment. The README file in the GitHub repository contains detailed information on how to set this up to reproduce the contents of the current manuscript, including a video.

³ Technically, this maximum entropy condition is satisfied because the $\text{beta}(1,1)$ distribution is uniform over its

93 we have a continuous scale with upper and lower bounds—and no other special conditions—the beta
94 distribution will in principle provide a very good approximation of the uncertainty of the scale.

95 Typically, the expected value (or mean) of the response variable is the central estimand
96 scholars want to estimate. A model should specify how this expected value depends on
97 explanatory variables through two main components: a linear predictor, which combines the
98 explanatory variables in a linear form ($a + b_1x_1 + b_2x_2$, etc.), and a link function, which connects
99 the expected value of the response variable to the linear predictor (e.g.,
100 $E[Y] = g(a + b_1x_1 + b_2x_2)$). In addition, a random component specifies the distribution of the
101 response variable around its expected value (such as Poisson or binomial distributions, which
102 belong to the exponential family) (Nelder & Wedderburn, 1972). Together, these components
103 provide a flexible framework for modeling data with different distributional properties.

104 The beta distribution is continuous and restricted to values between 0 and 1 (exclusive). Its
105 two parameters—commonly called shape1 (α) and shape2 (β)—govern the distribution’s location,
106 skewness, and spread. By adjusting these parameters, the distribution can take many functional
107 forms (e.g., it can be symmetric, skewed, U-shaped, or even approximately uniform; see Figure 1).

108 To illustrate, consider a test question worth seven points. Suppose a participant scores five
109 out of seven. The number of points received (5) can be treated as α , and the number of points
110 missed (2) as β . The resulting beta distribution would be skewed toward higher values, reflecting
111 a high performance (pink line in Figure 1; “beta(5, 2)”). Reversing these values would produce a
112 distribution skewed toward lower values, representing poorer performance (orange line in
113 Figure 1; “beta(2, 5)”).

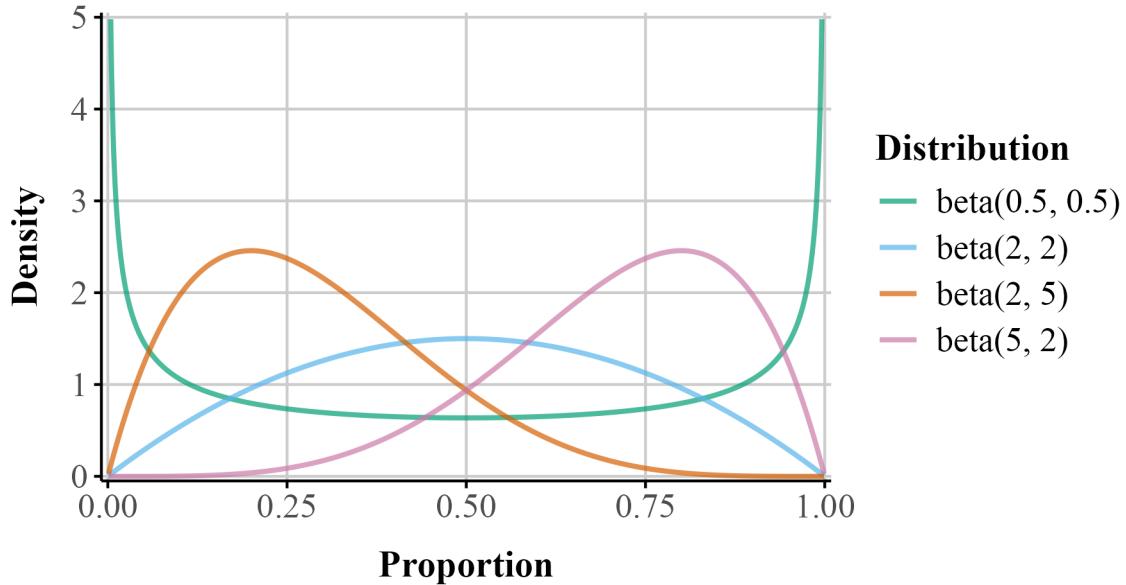
114 I Can’t Believe It’s Not beta

115 While the standard parameterization of the beta distribution uses α and β , a
116 reparameterization to a mean (μ) and precision (ϕ) is more useful for regression models. The
117 mean represents the expected value of the distribution, while the dispersion, which is inversely
118 related to variance, reflects how concentrated the distribution is around the mean, with higher

support. In addition, we assume that the scale has been re-scaled to the [0, 1] interval as we describe above.

Figure 1

beta distributions with different shape1 and shape2 parameters.



119 values indicating a narrower distribution and lower values indicating a wider one. The
 120 connections between the beta distribution's parameters are shown in Equation 1. Importantly, the
 121 variance depends on the average value of the response because uncertainty intervals need to adjust
 122 for how close the value of the response is to the boundary.

$$\text{Shape 1: } a = \mu\phi \quad \text{Mean: } \mu = \frac{a}{a+b} \quad (1)$$

$$\text{Shape 2: } b = (1 - \mu)\phi \quad \text{Precision: } \phi = a + b$$

$$\text{Variance: } var = \frac{\mu \cdot (1 - \mu)}{1 + \phi}$$

123 Thus, beta regression allows modeling both the mean and precision of the outcome
 124 distribution. To ensure that μ stays between 0 and 1, we apply a link function, which allows linear
 125 modeling of the mean on an unbounded scale. A common link-function choice is the logit, but
 126 other functions such as the probit or complementary log-log are possible.

127 The logit function, $\text{logit}(\mu) = \log\left(\frac{\mu}{1-\mu}\right)$ links the mean to log-odds which are unbounded,
 128 making linear modeling possible. The logit here no longer carries the same literal *odds*

129 interpretation because there are no corresponding counts of “successes” and “failures.” Instead,
130 the logit transform here simply maps the mean of the distribution to the real line. The inverse of
131 the logit, called the logistic function, maps the linear predictor η back to the original scale of the
132 data ($\mu = \frac{1}{1+e^{-\eta}}$). The coefficients describe how predictors shift the *average proportion* on the
133 logit scale. Similarly, the strictly positive dispersion parameter is usually modeled through a log
134 link function, ensuring it remains positive.

135 By accounting for the observations’ natural limits and non-constant variance, the beta
136 distribution is useful in psychology where outcomes like performance rates or response scales
137 frequently exhibit these features.

138 Bayesian Approach to Beta Regression

139 Beta regression models can be estimated using either frequentist or Bayesian methods. In
140 this paper, we adopt a Bayesian framework because it facilitates the estimation and interpretation
141 of more complex models (Gelman et al., 2013; Johnson et al., 2022; McElreath, 2020).
142 Additionally, the use of Bayesian statistics in psychology has been steadily growing (Pfadt et al.,
143 2025). In principle, frequentist methods like maximum likelihood can be framed as Bayesian
144 models with uninformative priors, and as a result, the modeling perspective we put forward in this
145 paper can apply to either approach. Nonetheless, we note that in non-linear and hierarchical
146 models, frequentist estimation may require additional adjustments such as bootstrapping to obtain
147 proper uncertainty intervals, whereas Bayesian modeling handles these extensions more naturally
148 via exploration of the full joint posterior distribution.⁴

149 There are several important differences between our Bayesian analysis and the frequentist

⁴ A common concern is that Bayesian methods are slower than frequentist ones. While this is true in general, modern Bayesian computation engines are efficient, and in explanatory modeling our priority is to specify appropriate estimands rather than to optimize for computation speed. Moreover, we use the {brms} package (Bürkner, 2017), which provides a high-level interface to Stan (Team, 2023) and uses familiar R formula syntax (similar to `lm()`), making advanced Bayesian modeling accessible to researchers with standard regression backgrounds. The package also supports parallelization, which substantially reduces computation time for large datasets.

150 methods readers may be more familiar with—most notably, the absence of t - and p -values. To
151 estimate models, the `{brms}` package uses Stan’s computational algorithms to draw random
152 samples from the posterior distribution, which represents uncertainty about the model parameters.
153 This posterior is conceptually analogous to a frequentist sampling distribution. By default,
154 Bayesian models run 4 chains with 2,000 iterations each.⁵ The first 1,000 iterations per chain are
155 warmup and are discarded. The remaining 1,000 iterations per chain are retained as posterior
156 draws, yielding 4,000 total post-warmup draws across all chains. From these draws, we can
157 compute the posterior mean (analogous to a frequentist point estimate) and the 95% credible
158 interval (Cr.I.), which is often compared to a confidence interval.

159 In addition, an important part of Bayesian analyses is prior specification. Priors encode
160 our assumptions about plausible parameter values before observing the data and allow the model
161 to regularize estimates, especially when data are sparse or parameters are weakly identified. To
162 help bridge the conceptual gap for users more familiar with frequentist models, we begin with the
163 default priors (flat/non-informative, or weakly informative in some cases) provided by `{brms}`.
164 These priors are intentionally non-informative, and in many applications produce results that
165 closely align with frequentist estimates, while still offering the flexibility and interpretive
166 advantages of a Bayesian framework. We strongly urge readers to consider prior specification
167 strongly in all their work.

168 To ease readers into Bayesian data analysis we provide a metric known as the *probability*
169 *of direction* (`pd`), which reflects the probability that a parameter is positive or negative. When a
170 uniform prior is used (all values equally likely in the prior), `pds` of 95%, 97.5%, 99.5%, and
171 99.95% corresponds approximately to two-sided p -values of .10, .05, .01, and .001 (i.e., $pd \approx 1 -$
172 $p/2$ for symmetric posteriors with weak/flat priors) (see Figure 2 for an illustrative comparison).
173 For directional hypotheses, the `pd` can be interpreted as roughly equivalent to one minus the

⁵ The Hamiltonian Monte Carlo sampler employed by Stan, which we also use in this paper, can converge with significantly fewer iterations, though rapid convergence depends on model complexity, which is why we use a more conservative standard in this paper.

¹⁷⁴ *p*-value (Marsman & Wagenmakers, 2016).

¹⁷⁵ For reasons of space, we refer readers unfamiliar to Bayesian data analysis to several
¹⁷⁶ existing books on the topic (Gelman et al., 2013; Kruschke, 2015; McElreath, 2020). In addition,
¹⁷⁷ we assume readers are familiar with R, but those in need of a refresher should find Wickham et al.
¹⁷⁸ (2023) useful.

¹⁷⁹ **Beta Regression Tutorial**

¹⁸⁰ **Example Data**

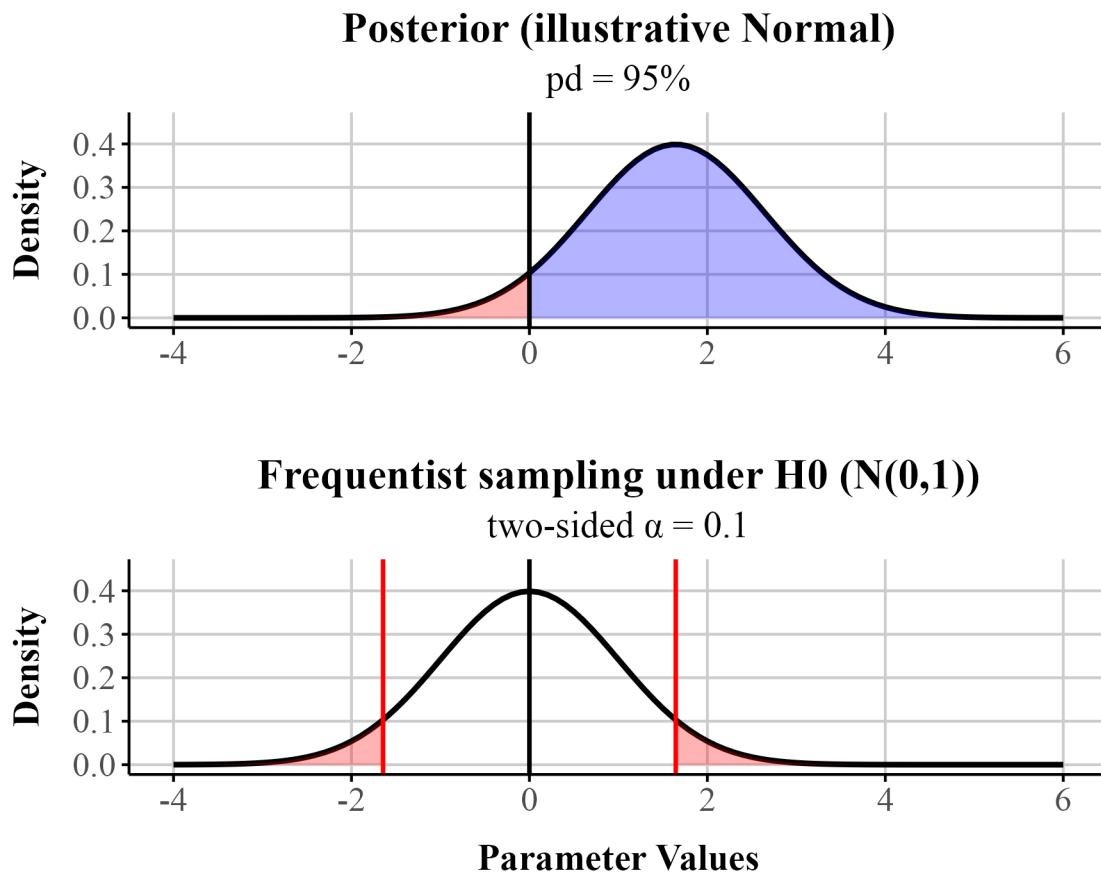
¹⁸¹ Throughout this tutorial, we analyze data from a memory experiment examining whether
¹⁸² the fluency of an instructor's delivery affects recall performance (Wilford et al., 2020, Experiment
¹⁸³ 1A). Instructor fluency—marked by expressive gestures, dynamic vocal tone, and confident
¹⁸⁴ pacing—has been shown to influence students' perceptions of learning, often leading learners to
¹⁸⁵ rate fluent instructors more favorably (Carpenter et al., 2013). However, previous research
¹⁸⁶ suggests that these impressions do not reliably translate into improved memory performance (e.g.,
¹⁸⁷ Carpenter et al., 2013; Toftness et al., 2017; Witherby & Carpenter, 2022). In contrast, Wilford et
¹⁸⁸ al. (2020) found that participants actually recalled more information after watching a fluent
¹⁸⁹ instructor compared to a disfluent one. This surprising finding makes the dataset a compelling
¹⁹⁰ case study for analyzing proportion data, as recall was scored out of 10 possible idea units per
¹⁹¹ video.

¹⁹² In Experiment 1A, ninety-six participants watched two short instructional videos, each
¹⁹³ delivered either fluently or disfluently. Fluent videos featured instructors with smooth delivery
¹⁹⁴ and natural pacing, while disfluent videos included hesitations, monotone speech, and awkward
¹⁹⁵ pauses. After a distractor task, participants completed a free recall test, writing down as much
¹⁹⁶ content as they could remember from each video within a three-minute window. Their recall was
¹⁹⁷ then scored for the number of idea units correctly remembered.

¹⁹⁸ Our primary outcome variable is the proportion of idea units recalled on the final test,
¹⁹⁹ calculated by dividing the number of correct units by 10. We show a sample of these data in
²⁰⁰ Table 1. The dataset can be downloaded from GitHub (Listing 1). Because this is a bounded

Figure 2

A Bayesian posterior distribution (assuming a uniform prior) centered at a point estimate chosen so that the probability of direction (pd) equals 0.95, and a frequentist sampling distribution (under the null; centered at 0). In the Bayesian posterior distribution, the blue area represents the pd, and the red area represents the remaining $1 - pd$ of the distribution. In the frequentist sampling distribution, the red tail areas represent the rejection region at $\alpha = 0.10$. In this example, the posterior mean lies exactly at the $1 - \frac{\alpha}{2}$ quantile of the null sampling distribution. For symmetric posteriors with flat priors, the pd is numerically equivalent to the one-sided p-value.



Listing 1 Data needed to run examples

```
# get data here from project folder
fluency_data <- read_csv(here::here("manuscript", "data", "fluency_data.csv"))
```

201 continuous variable (i.e., it ranges from 0 to 1), it violates the assumptions of typical linear
 202 regression models that assume normally distributed residual errors. Despite this, it remains
 203 common in psychological research to analyze proportion data using models that assume
 204 normality. In what follows, we reproduce Wilford et al. (2020)'s analysis and then re-analyze the
 205 data using beta regression and highlight how it can improve our inferences.

Table 1

Four observations from Wilford et al. (2020). Accuracy refers to the proportion of correctly recalled idea units.

| Participant | Fluency | Accuracy |
|-------------|-----------|----------|
| 64 | Disfluent | 0.10 |
| 30 | Fluent | 0.60 |
| 12 | Fluent | 0.10 |
| 37 | Fluent | 0.35 |

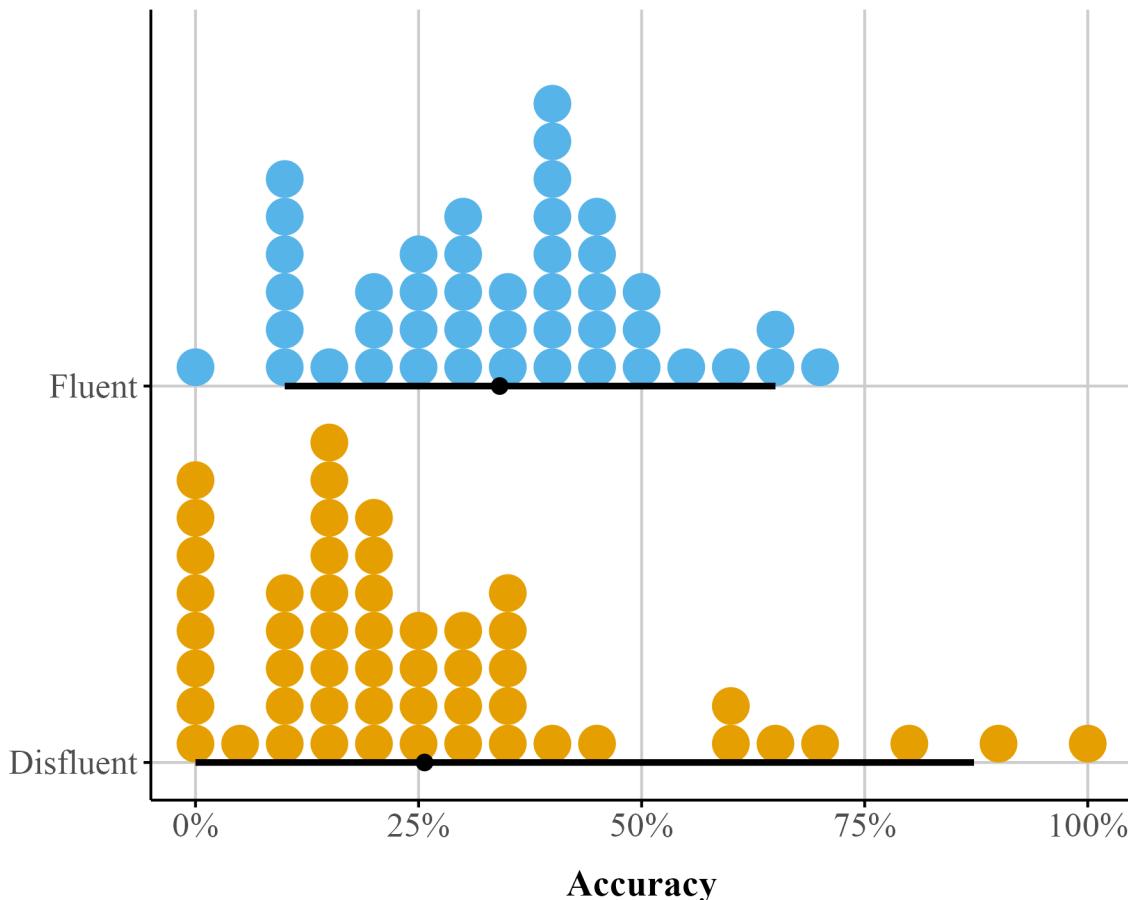
206 Reanalysis of Wilford et al. Experiment 1A

207 In their original analysis of Experiment 1A, Wilford et al. (2020) compared memory
 208 performance between fluent and disfluent instructor conditions using a traditional
 209 independent-samples t-test on mean accuracy for 96 participants. They found that participants
 210 who watched the fluent instructor recalled significantly more idea units than those who viewed the
 211 disfluent version (see Figure 3).

212 We first replicate this analysis in a regression framework using {brms}. We model final
 213 test mean accuracy—the proportion of correctly recalled idea units across the videos—as the

Figure 3

Dot plot depicting accuracy distributions for the Fluent and Disfluent conditions. Each condition shows individual data points and summary statistics (mean and 95% percentile interval) to illustrate variability and central tendency.



214 dependent variable. Our predictor is instructor fluency, with two levels: Fluent and Disfluent. We
 215 use treatment (dummy) coding, which is the default in R. This coding scheme sets the first level of
 216 a factor (in alphabetical order) as the reference level. In this case, Disfluent is the reference, and
 217 the coefficient for Fluent reflects the contrast between fluent and disfluent instructor conditions.

218 **Regression Analysis**

219 We first start by loading the `{brms}` (Bürkner, 2017) and `{cmdstanr}` (Gabry et al., 2024)
 220 packages (Listing 2). We use the cmdstanr backend for Stan (Team, 2023) because it's faster than

221 the default used to run models (i.e., `rstan`)⁶ though all of these models can also be fit with `brms`
222 defaults.

Listing 2 Load the `{brms}` and `{cmdstanr}` packages

```
library(brms)
library(cmdstanr)
```

Listing 3 Fitting a gaussian model with `brm()`.

```
bayes_reg_model <- brm(
  Accuracy ~ Fluency,
  data = fluency_data,
  family = gaussian(),
  file = here::here("manuscript", "models", "model_reg_bayes")
)
```

223 We fit the model using the `brm()` function from the `{brms}` package (Listing 3). Although
224 not shown here, we ran the models using four chains (the default), executed in parallel across four
225 cores. When the model is run in Listing 3, the model summary output will appear in the R
226 console. The output from `bayes_reg_model` shows each parameter's posterior summary: The
227 posterior distribution's mean and standard deviation (analogous to the frequentist standard error)
228 and its 95% credible interval, which indicate the 95% of the most credible parameter values. In
229 `{brms}`, the reported Cr.I is an equal-tailed interval, meaning that the probability mass excluded
230 from the interval is split equally between the lower and upper tails. Additionally, the output
231 indicates numerical estimates of the sampling algorithm's performance: Rhat should be close to
232 one, and the ESS (effective sample size) metrics should be as large as possible given the number

⁶ In order to use the `cmdstanr` backend you will need to first install the package (see
<https://mc-stan.org/cmdstanr/>) and also run `cmdstanr::install_cmdstan()` if you have not done so
already.

233 of iterations specified (default is 4000). Generally, ESS ≥ 1000 is recommended (Bürkner,
 234 2017). For the models we present in this paper, convergence is trivial with standard linear models,
 235 though we note that these metrics are still important to pay attention to in case of model misfit.

236 Our main question of interest is: what is the causal effect of instructor fluency on final test
 237 performance? In order to answer this question, we will have to look at the output summary
 238 produced by Listing 3 (also see Table 8 under Bayesian LM). the Intercept refers to the
 239 posterior mean accuracy in the disfluent condition, $M = 0.256$, as fluency was dummy-coded.
 240 The fluency coefficient (FluencyFluent) reflects the mean posterior difference in recall accuracy
 241 between the fluent and disfluent conditions: $b = 0.085$. The 95% Cr.I for this estimate spans from
 242 0 to 0.167. These values are shown in the “95% Cr.I” columns of the output. These results closely
 243 mirror the findings reported by Wilford et al. (2020) (Experiment 1A).

244 Family: gaussian
 245 Links: mu = identity
 246 Formula: Accuracy ~ Fluency
 247 Data: fluency_data (Number of observations: 96)

248
 249 Regression Coefficients:

| | Estimate | Est.Error | l-95% | CI | u-95% | CI | Rhat | Bulk_ESS | Tail_ESS |
|---------------|----------|-----------|-------|------|-------|------|------|----------|----------|
| Intercept | 0.26 | 0.03 | 0.20 | 0.31 | 1.00 | 4190 | 2492 | | |
| FluencyFluent | 0.08 | 0.04 | 0.00 | 0.17 | 1.00 | 4386 | 3152 | | |

253
 254 Further Distributional Parameters:

| | Estimate | Est.Error | l-95% | CI | u-95% | CI | Rhat | Bulk_ESS | Tail_ESS |
|-------|----------|-----------|-------|------|-------|------|------|----------|----------|
| sigma | 0.21 | 0.02 | 0.18 | 0.24 | 1.00 | 4259 | 2944 | | |

257 The output also includes the effective sample size (ESS) and R (R-hat) values, both of
 258 which fall within acceptable ranges, indicating good model convergence. Throughout the tutorial,
 259 we focus primarily on posterior mean estimates and their 95% credible intervals. In addition, we

260 report the pd measure in the main summary table (Table 8), provided by the {bayestestR} package
 261 (Makowski, Ben-Shachar, Chen, et al., 2019; Makowski, Ben-Shachar, & Lüdecke, 2019). This
 262 measure offers an intuitive parallel to *p*-values, which many readers may find familiar. For
 263 example, the fluency effect has a pd of .977, indicating a high probability that the effect is positive
 264 rather than negative.

265 Importantly, pd does not indicate whether an effect is meaningfully different from a point
 266 value—it only reflects the proportion of the posterior in one direction. To address questions of
 267 practical significance, readers can consider the Region of Practical Equivalence (ROPE) with the
 268 Cr.Is (Kruschke, 2015). Unlike a hypothesis test of a point null (e.g., exactly zero), the ROPE
 269 defines a range of values that are deemed too small to be of substantive importance. As a rule of
 270 thumb (see Kruschke, 2018), if more than 95% of the posterior distribution lies inside the ROPE,
 271 the effect can be considered practically equivalent to that negligible range. If less than 5% lies
 272 inside, the effect can be considered meaningfully different. Intermediate cases are typically
 273 labeled undecided.

274 The `rope()` function in the {bayestestR} package computes the proportion of the posterior
 275 within the ROPE to facilitate this evaluation. By default, from bayesian models fit via {brms}, the
 276 package determines a ROPE based on the data (roughly reflecting “negligible” effects), but these
 277 defaults should be used cautiously. The choice of ROPE should always be guided by theoretical
 278 considerations, previous research, and the substantive context of the study. In Listing 4, we show
 279 how to compute this using {bayestestR}. Running the function with default settings suggests that
 280 less than 5% lies within the default ROPE (indicating the effect is larger than .02) (see Figure 4).

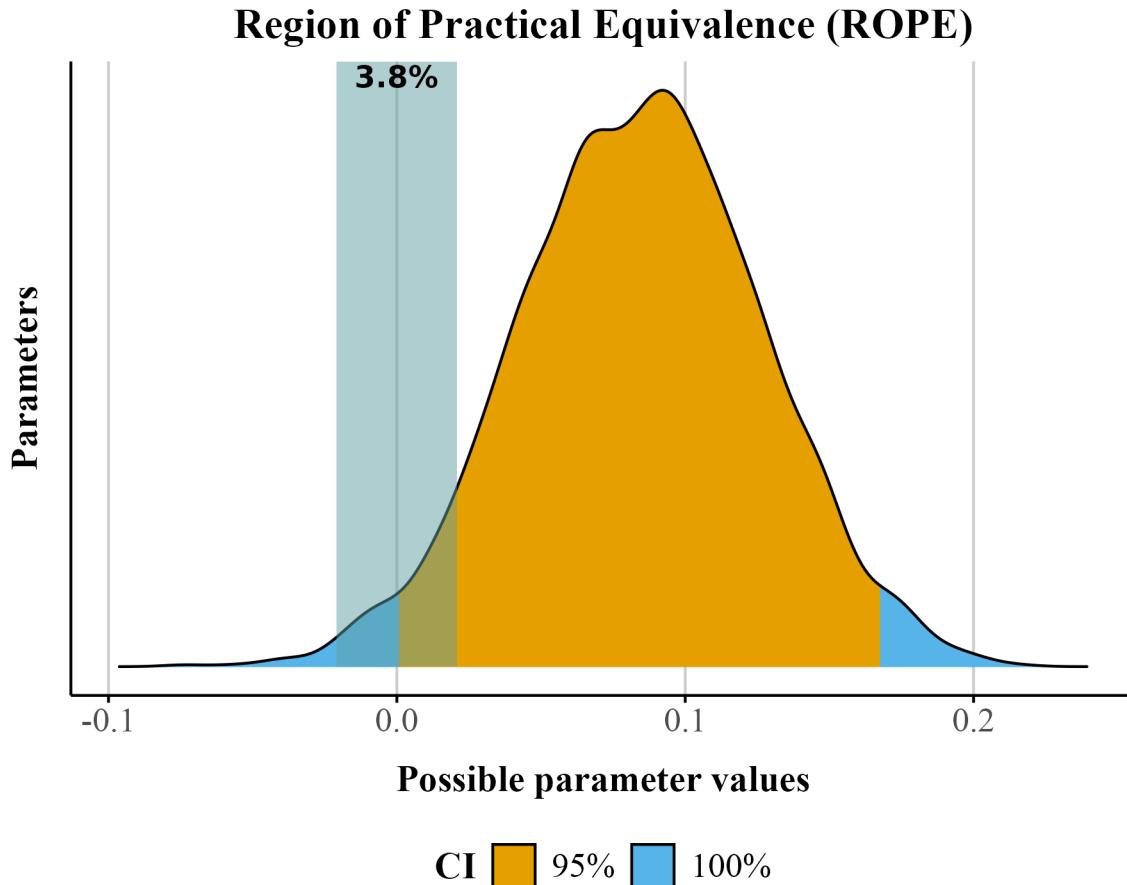
Listing 4 Getting ROPE from `bayes_reg_model` obect using `rope` function from {bayestestR}

```
brms_rope <- bayestestR::rope(bayes_reg_model, ci = .95, ci_method = "ETI")
```

281 **Assumptions.** Wilford et al. (2020) observed that instructor fluency impacts actual
 282 learning, using a standard *t*-test on the mean accuracy. But recall this approach assumes normality
 283 of residuals and homoskedasticity. These assumptions are unrealistic when the response values

Figure 4

Posterior distribution for the fluency effect showing the ROPE(shaded area) with 95% credible interval (orange) and 100% credible interval (blue). The percentage indicates the proportion of the posterior within the ROPE.



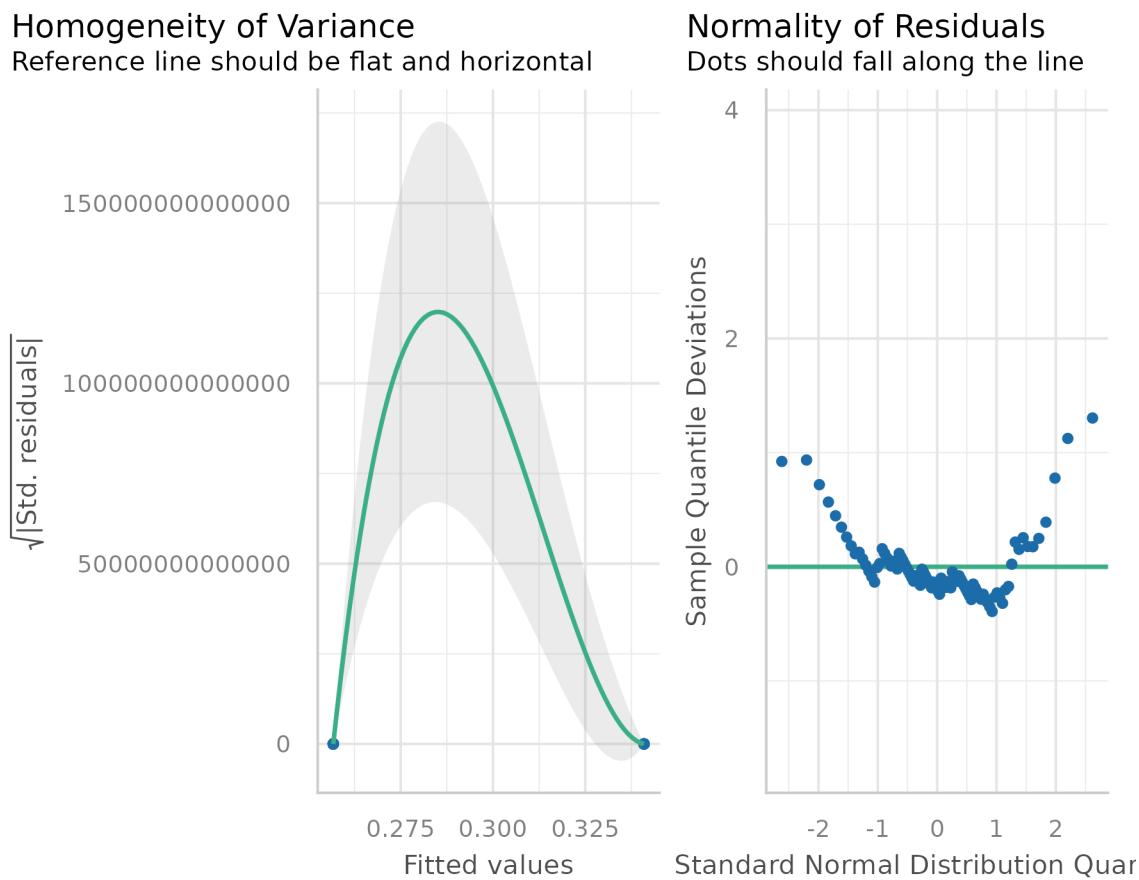
²⁸⁴ approach the scale boundaries (Sladekova & Field, 2024). Does the data we have meet those
²⁸⁵ assumptions? We can use the function `check_model()` from `{easystats}` (Lüdecke et al., 2022) to
²⁸⁶ check our assumptions easily. The code in Listing 5 produces Figure 5. We can see some issues
²⁸⁷ with our model. Specifically, there appears to be violations of constant variance across the values
²⁸⁸ of the scale (homoskedasticity). In plain terms, this type of model mis-specification means that a
²⁸⁹ standard OLS model can predict non-sensical values outside the bounds of the scale.

Listing 5 Checking assumptions with the `check_model()` from `{easystats}` package.

```
check_model(bayes_reg_model, check = c("homogeneity", "qq"))
```

Figure 5

Two assumption checks for our OLS model: Normality (right) and Homoskedasticity (left)



290 We can also examine how well the data fits the model by performing a posterior predictive

291 check using the `pp_check()` function from `{brms}`. A posterior predictive check involves looking
 292 at multiple model-predicted values and plotting them against the observed data. Ideally, the
 293 predicted values (the light blue lines) should show reasonable resemblance with the observed data
 294 (dark blue line). In our example (see Figure 12 (A)) the model-predicted density is slightly too
 295 peaked and narrow compared to the data. In addition, some of the predicted accuracy values are

296 negative.

297 **Distributional Regression - Beta**

298 It is important to note that there are several justifiable approaches for addressing the
299 distributional issues observed in the data. For instance, one could analyze median accuracy
300 instead of the mean, use some type of robust estimator for heterogeneity, or apply non-parametric
301 methods to relax some of the model assumptions. Alternatively, we can address these issues
302 directly by fitting distributional models (Kneib et al., 2023; Kruschke, 2013). A key advantage of
303 distributional models is that they are not limited to modeling only the mean or median of the
304 outcome, but can also model parameters such as the variance (or other shape parameters) as
305 functions of predictors. This allows examining how instructor fluency may influence not only
306 average performance, but also the variability in performance across students. If we wanted to keep
307 our mean accuracy variable and continue to use a Gaussian model, we could use a distributional
308 approach and model the effect of fluency on σ .

309 Given the outcome variable is proportional, another solution would be to run a beta
310 regression model. Again, we can create the beta regression model in {brms}. In {brms}, we
311 model each parameter independently. Recall from the introduction that in a beta model we model
312 two parameters— μ and ϕ . Again we do this by using the bf() function from {brms} (Listing 6).
313 We specify two formulas, one for μ and one for ϕ and store it in the model_beta_bayes object
314 below. In the below bf() call, we are modeling accuracy as a function of fluency only for the μ
315 parameter. For the ϕ parameter, we are only modeling the intercept value. This is saying
316 dispersion does not change as a function of fluency.

317 To run our beta regression model, we need to exclude 0s and 1s in our data set. If we try to
318 run a model with our data data_fluency we get an error: Error: Family 'beta' requires
319 response greater than 0. This is because the beta distribution only supports observations in
320 the 0 to 1 interval *excluding exact 0s and 1s*. We need make sure there are no 0s and 1s in our
321 dataset.

322 The dataset contains nine 0s and one 1. One approach is to nudge our 0s towards .01 and

323 our 1s to .99, or apply a special formula (Smithson & Verkuilen, 2006) so values fall within the [0,
324 1] interval. We implore readers not to engage in this practice. Kubinec (2022) showed that this
325 practice can result in serious distortion of the outcome as the sample size grows larger, resulting
326 in ever smaller values that are “nudged”. Because the beta distribution is a non-linear model of
327 the outcome, values that are very close to the boundary, such as 0.00001 or 0.99999, will be
328 highly influential outliers. To run this beta model we will remove the 0s and 1s, and later in this
329 article we will show how to jointly model these scale end points with the rest of the data. The
330 model from Listing 6 uses a transformed `data_fluency` object (called `data_beta`) where 0s and
331 1s are removed. When we run this code we should not get an error.

332 **Model Parameters.** In Table 8, under the beta regression column, the coefficient with b_{fluency}
333 represents how fluency of instructor influences the μ parameter estimates (which is the mean of
334 the distribution here). These coefficients are linear on the logit-scale, but not on the raw accuracy
335 scale. The intercept term ($b_{\text{Intercept}}$) represents the log odds of the mean on accuracy for the
336 fluent instructor. Log odds that are negative indicate that it is more likely a “success” (like getting
337 the correct answer) will not happen than that it will happen. Similarly, regression coefficients in
338 log odds forms that are negative indicate that an increase in that predictor leads to a decrease in
339 the predicted probability of a “success”.

340 The other component we need to pay attention to is the dispersion or precision parameter
341 coefficients labeled as ϕ in Table 8. The dispersion (ϕ) parameter tells us how precise our
342 estimate is. Specifically, ϕ in beta regression tells us about the variability of the response variable
343 around its mean. Specifically, a higher dispersion parameter indicates a narrower distribution,
344 reflecting less variability. Conversely, a lower dispersion parameter suggests a wider distribution,
345 reflecting greater variability. The main difference between a dispersion parameter and the
346 variance is that the dispersion has a different interpretation depending on the value of the
347 outcome, as we show below. The best way to understand dispersion is to examine visual changes
348 in the distribution as the dispersion increases or decreases.

349 Understanding the dispersion parameter helps us gauge the precision of our predictions

Listing 6 Fitting a beta model without 0s and 1s in brm().

```
# set up model formual

model_beta_bayes <- bf(
  Accuracy ~ Fluency, # fit mu model
  phi ~ 1 # fit phi model
)

# transform 0 to 0.1 and 1 to .99

data_beta <- fluency_data |>
  filter(
    Accuracy != 0,
    Accuracy != 1
  )

beta_brms <- brm(
  model_beta_bayes,
  data = data_beta,
  family = Beta(),
  file = here::here("manuscript", "models", "model_beta_bayes_reg_01")
)
```

350 and the consistency of the response variable. In `beta_brms` we only modeled the dispersion of
 351 the intercept. When ϕ is not specified, the intercept is modeled by default (see Table 8). It
 352 represents the overall dispersion in the outcome across all conditions. Instead, we can model
 353 different dispersions across levels of the Fluency factor. To do so, we add `Fluency` to the `phi`
 354 model in `bf()`. We model the precision (`phi`) of the `Fluency` factor by using a `~` and adding
 355 factors of interest to the right of it (Listing 7).

Listing 7 Fitting beta model with dispersion.

```
model_beta_bayes_disp <- bf(
  Accuracy ~ Fluency, # Model of the mean
  phi ~ Fluency # Model of the precision
)

beta_brms_dis <- brm(
  model_beta_bayes_disp,
  data = data_beta,
  family = Beta(),
  file = here::here("manuscript", "models", "model_beta_bayes_dis_run01")
)
```

356 Table 8 displays the model summary with the precision parameter labeled as

357 phi_Fluency. Since ϕ is modeled on the log scale, the coefficients represent changes in $\log\phi$
 358 rather than ϕ itself. To see the effect in the original units, we convert the values back by
 359 exponentiating. Thus, the effect of the Fluent condition can be understood by comparing the
 360 exponentiated predicted ϕ in the Fluent condition to that in the baseline condition.

361 The ϕ parameters are estimated on the log scale. The term $\beta_{\text{Intercept}}^{(\phi)}$ represents the
 362 log-precision for the reference (disfluent) condition. The coefficient $\beta_{\text{FluencyFluent}}^{(\phi)}$ represents the
 363 change in log-precision when moving from the disfluent to the fluent condition.

364 To obtain precision on the original scale, we exponentiate the linear predictor:

$$\phi_{\text{disfluent}} = \exp(\beta_{\text{Intercept}}^{(\phi)}), \quad \phi_{\text{fluent}} = \exp(\beta_{\text{Intercept}}^{(\phi)} + \beta_{\text{Fluency}}^{(\phi)}).$$

365 The coefficient $\beta_{\text{Fluency}}^{(\phi)}$ therefore describes a *multiplicative* change in precision.

366 Specifically,

$$\frac{\phi_{\text{fluent}}}{\phi_{\text{disfluent}}} = \exp(\beta_{\text{Fluency}}^{(\phi)}).$$

Because the 95% credible interval for $\beta_{\text{Fluency}}^{(\phi)}$ does not include zero, we infer that there is

a credible difference in precision between the fluent and disfluent conditions.

It is important to note that these estimates are not the same as the marginal effects we

discussed earlier. Changes in dispersion affect the spread or variability of the response

distribution without necessarily altering its mean. This makes dispersion particularly relevant for

research questions that focus on features of the distribution beyond the average—such as how

concentrated responses are. For instance, high dispersion might indicate that individuals cluster at

the extremes (e.g., very high or very low ratings), suggesting clustering in the outcome.

A critical assumption of the linear model is homoscedasticity, which means constant

variance of the errors. With beta regression model we can include a dispersion parameter for

Fluency. Properly accounting for dispersion is crucial because it impacts the precision of our mean

estimates and, consequently, the substantive inferences we might make about the coefficients. The

inclusion of dispersion in the model increased the uncertainty of the μ coefficient (see Figure 6).

This highlights the potential utility of an approach like beta regression over a traditional approach

as beta regression can explicitly model dispersion and address issues of heteroscedasticity.

While it is advisable to model precision, if there is uncertainty about the best model, a

relatively agnostic approach would be to compare models, for example with leave one out (loo)

cross validation, to examine if a dispersion parameter should be considered in our model.⁷

385 Predicted Probabilities

Parameter estimates can be difficult to interpret, and researchers can instead discuss effects

on the actual outcome scale (in this case the 0-1 scale). The logit link allows us to transform back

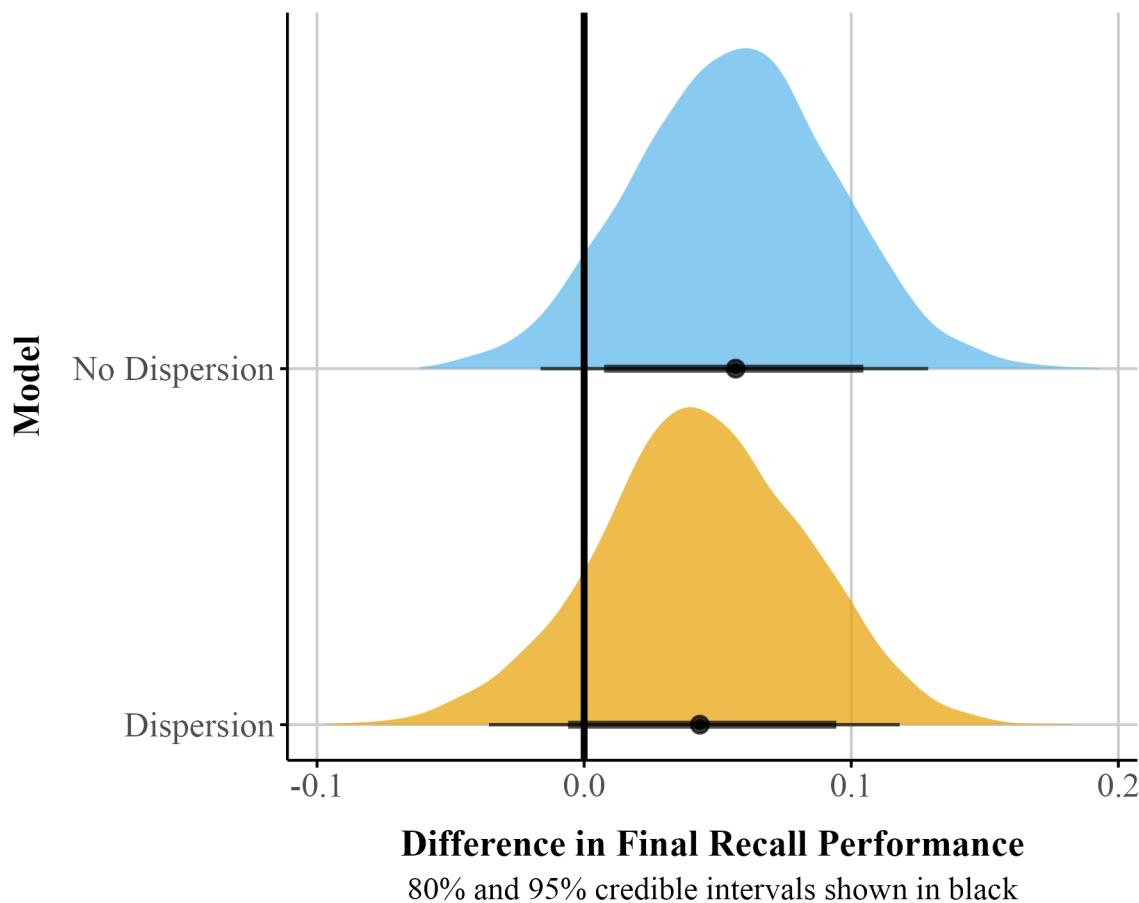
and forth between the scale of a linear model and the nonlinear scale of the outcome, which is

bounded by 0 and 1. By using the inverse of the logit, we can easily transform our linear

⁷ The model fit statistic LOO-CV can be compared for any set of fitted brms models with the function loo().

Figure 6

Comparison of posterior distributions for the risk difference in fluency: Simple model (no dispersion for Fluency) vs. complex model with dispersion



390 coefficients to obtain average effects on the scale of the proportions or percentages, which is
 391 usually easier to interpret. In a simple case, we can do this manually, but when there are many
 392 factors in your model this can be quite complex.

393 In our example, we can use the `plogis()` function in base R to convert estimates from the
 394 logit scale to the probability scale. The intercept of our model is -0.918, which reflects the logit of
 395 the mean accuracy in the disfluent condition. If the estimated difference between the fluent and
 396 disfluent conditions is 0.24 on the logit scale, we first add this value to the intercept value (-0.918)
 397 to get the logit for the fluent condition: $-0.83 + 0.20 = -0.63$. We then use `plogis()` to
 398 convert both logit values to probabilities (Fluent = 35%, Disfluent = 30%).

399 With single coefficients this calculation is trivial, but in more complex models with
 400 interactions, it can be quite cumbersome. To help us extract predictions from our model and
 401 visualize them we will use a package called `{marginaleffects}` (Arel-Bundock et al., 2024) (see
 402 Listing 8). To get the proportions for each of our categorical predictors on the μ parameter we can
 403 use the function from the package called `avg_predictions()`. These are displayed in Table 2.
 404 These probabilities match what we calculated above.

Listing 8 Load the `{marginaleffects}` package.

```
library(marginaleffects)

options(marginaleffects_posterior_center = mean) # make sure returns mean
```

Listing 9 Predictions from the beta model for each level of Fluency (μ).

```
avg_predictions(
  beta_brms_dis,
  # need to specify the levels of the categorical predictor
  variables = "Fluency"
)
```

Table 2

Predicted probabilities for fluency factor (μ).

| Fluency | Mean | 95% Cr.I |
|-----------|-------|----------------|
| Disfluent | 0.304 | [0.249, 0.363] |
| Fluent | 0.348 | [0.301, 0.398] |

405 For the Fluency factor, we can interpret the estimates under the Mean column as a
 406 proportion or percentage. That is, participants who watched the fluent instructor scored on
 407 average 35% on the final exam compared to 30% for those who watched the disfluent instructor.

408 We can also visualize these from `{marginaleffects}` using the `plot_predictions()` function
 409 (see Listing 10).

Listing 10 Plot predicted probabilities using `plot_predictions()` from `{marginaleffects}`

```
beta_plot <- plot_predictions(beta_brms_dis, by = "Fluency")
```

410 The `plot_predictions()` function will only display the point estimate with the 95%
 411 credible interval. However, Bayesian estimation methods generate distributions for each
 412 parameter. This approach allows visualizing full uncertainty estimates beyond points and
 413 intervals. Using the `{marginaleffects}` package, we can obtain samples from the posterior
 414 distribution with the `posterior_draws()` function (see Listing 11). We can then plot these
 415 results to illustrate the range of plausible values for our estimates at different levels of uncertainty
 416 (see Figure 7).

Listing 11 Extracting posterior draws from the beta regression model.

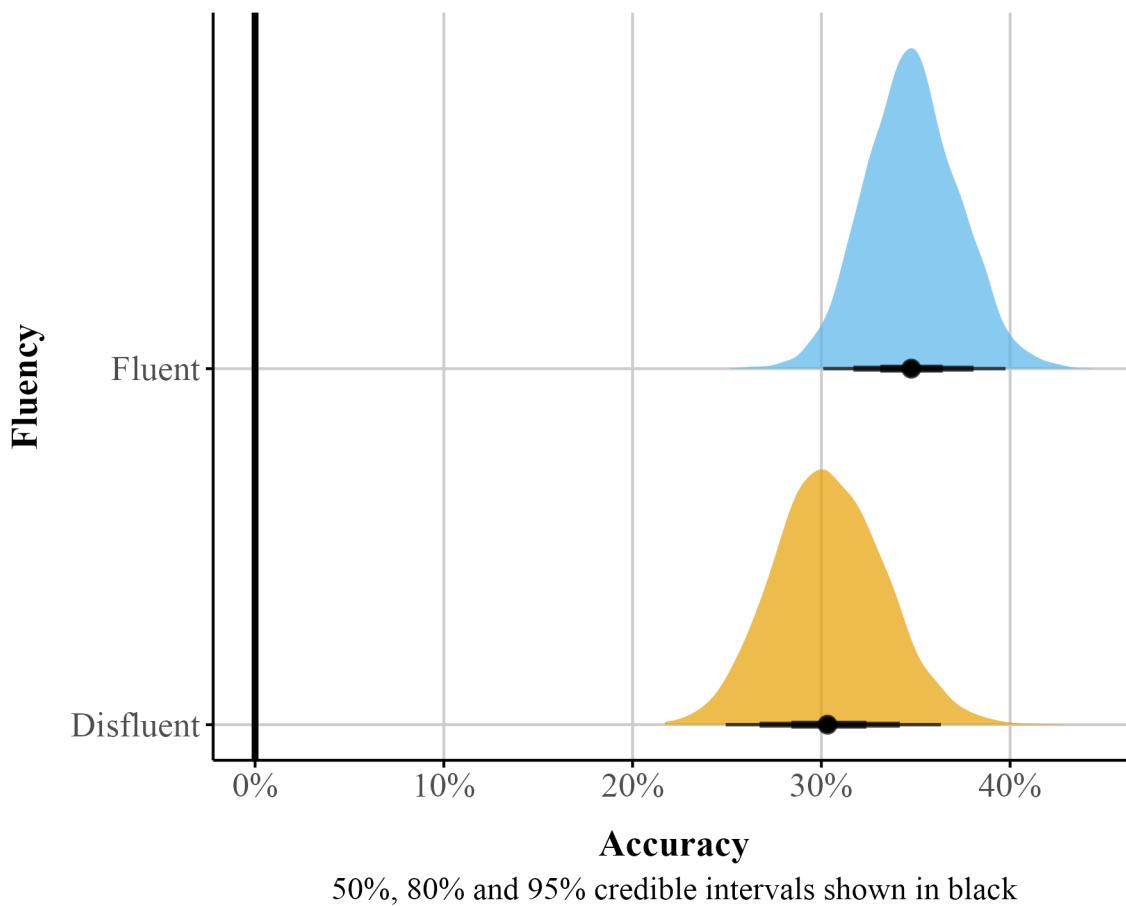
```
# Add a model identifier to each dataset
pred_draws_beta <- avg_predictions(beta_brms_dis, variables = "Fluency") |>
  posterior_draws()
```

417 ***Marginal Effects***

418 Marginal effects offer an interpretable way to quantify how changes in a predictor
 419 influence an outcome, while holding other factors constant in a specific manner. In recent years,
 420 there has been a thrust to move away from reporting regression coefficients alone, focusing instead
 421 on estimates that are easier to interpret and communicate—particularly in non-linear models
 422 (McCabe et al., 2021; Rohrer & Arel-Bundock, 2025). Technically, marginal effects are computed
 423 as partial derivatives for continuous variables or as finite differences for categorical (and
 424 sometimes continuous) predictors, depending on the structure of the data and the research
 425 question. Substantively, these procedures translate raw regression coefficients into quantities that

Figure 7

Predicted probability posterior distributions by instructor fluency



⁴²⁶ reflect changes in the bounded outcome—for example, an $x\%$ change in the value of a proportion.

⁴²⁷ There are several types of marginal effects, and their computation can vary across software
⁴²⁸ packages. For example, the popular `{emmeans}` package (Lenth, 2025) computes marginal effects
⁴²⁹ by holding all predictors at their means (marginal effects at the mean; MEM). In this tutorial, we
⁴³⁰ use the `{marginaleffects}` package (Arel-Bundock et al., 2024), which by default computes
⁴³¹ average marginal effects (AMEs). AMEs are based on counterfactual predictions: the dataset is
⁴³² conceptually replicated across all unique values of the predictor of interest, predictions are
⁴³³ generated for each row under each counterfactual scenario, and the resulting differences are then
⁴³⁴ averaged. This approach maintains a strong connection to the observed data—because predictions
⁴³⁵ are made using each participant’s actual values—while providing a clear and interpretable

436 summary of the effect of interest.

437 One practical use of AMEs is to estimate the average difference between two groups or
438 conditions which corresponds to the average treatment effect (ATE). Using the
439 `avg_comparisons()` function in the `{marginaleffects}` package (Listing 12), we can compute this
440 quantity directly. By default, the function returns the discrete difference between groups. When
441 we take the difference in proportions between two groups it is called the risk difference.
442 Depending on the audience and modeling goals, the function can also produce alternative effect
443 size metrics, such as odds ratios or risk ratios. This flexibility makes it a powerful approach for
444 summarizing and communicating regression results.

Listing 12 Calculating the difference between probabilities (μ) for fluency with `avg_comparisons()`

```
# get risk difference by default
beta_avg_comp <- avg_comparisons(
  beta_brmis_dis,
  variables = "Fluency", # counterfactual difference
  comparison = "difference"
)
```

Table 3

Fluency difference (μ)

| Contrast | Mean | 95% Cr.I | pd |
|--------------------|-------|-----------------|------|
| Fluent - Disfluent | 0.044 | [-0.036, 0.118] | 0.87 |

445 **μ Difference.** Table 3 presents the estimated difference for the Fluency factor (Mean
446 column). The difference between the fluent and disfluent conditions is 0.06, indicating that
447 participants who watched a fluent instructor scored, on average, 6% higher on the final recall test

448 than those who watched a disfluent instructor. However, the 95% credible interval includes 0
 449 among the most credible values, suggesting we cannot rule out the possibility of a null or weakly
 450 negative effect. The pd also indicates the effect is non-significant ($p > .05$). In the basic regression
 451 model above, the ROPE value was extracted using the `{bayestestR}` package.

Listing 13 Extracting ROPE using `{marginaleffects}` for μ difference

```
# get risk difference by default

beta_avg_comp_rope <- avg_comparisons(
  beta_brms_dis,
  variables = "Fluency", # counterfactual difference
  comparison = "difference",
  equivalence=c(-0.02, .02)
)
```

452 ,The ROPE can also be obtained using the `{marginaleffects}` package by specifying the
 453 equivalence bounds in `avg_comparisons()`. Running the code in Listing 13 returns a ROPE
 454 probability of 23%, indicating that there is insufficient evidence to conclude that the effect
 455 exceeds 2% or is practically equivalent.

456 **ϕ difference.** In addition, with `{marginaleffects}`, we can get the actual precision
 457 difference between the two groups on ϕ using similar code to above by setting dpar to “phi”
 458 (Listing 14).

459 **Effect Size.** In psychology, it is common to report effect size measures like Cohen’s d
 460 (Cohen, 1977). When working with proportions we can calculate something similar called
 461 Cohen’s h . Taking our proportions, we can use the below equation (Equation 2) to calculate
 462 Cohen’s h along with the 95% Cr.I around it. Using this metric we see the effect size is small
 463 (0.107), 95% credible interval [-0.002, 0.361].

$$h = 2 \cdot (\arcsin(\sqrt{p_1}) - \arcsin(\sqrt{p_2})) \quad (2)$$

Listing 14 Calculating ϕ difference with avg_comparisons()

```
# get risk difference by default

beta_avg_phi <- avg_comparisons(
  beta_brms_dis,
  variables = "Fluency",
  dpar = "phi",
  comparison = "difference"
)
```

464 **Posterior Predictive Check**

465 Figure 12 (B) shows the predictive check for our beta model. The model's predictions
466 generally conform to the data as the predictions are now between constrained to the 0-1 interval.
467 However, the model's predictive performance can be improved the bounds of the scale are
468 explicitly specified.

469 **Zero-Inflated beta (ZIB) Regression**

470 A limitation of the beta regression model is that it can only accommodate values strictly
471 between 0 and 1—a probability cannot take on values of 0 (the event will not occur with certainty)
472 or 1 (the event will occur with certainty). In our dataset, we observed 9 rows where Accuracy
473 equals zero. To fit a beta regression model, we removed these values, but we have left out
474 potentially valuable information from our model—especially if the end points of the scale are
475 distinctive in some way. In our case, these 0s may be structural—that is, they represent real,
476 systematic instances where participants failed to answer correctly (rather than random noise or
477 measurement error). For example, the fluency of the instructor might be a key factor in predicting
478 these zero responses. We will discuss two approaches for jointly modeling these end points with
479 the continuous data. First, we can use a zero-inflated beta (ZIB) model. This model still estimates
480 the mean (μ) and precision (ϕ) of the beta distribution for values between 0 and 1, but it also

481 includes an additional parameter, α , which captures the probability of observing structural 0s.

482 The zero-inflated beta models a mixture of the data-generating process. The α parameter
483 uses a logistic regression to model whether the data is 0 or not. Substantively, this could be a
484 useful model when we think that 0s come from a process that is relatively distinct from the data
485 that is greater than 0. For example, if we had a dataset with proportion of looks or eye fixations to
486 certain areas on marketing materials, we might want a separate model for those that do not look at
487 certain areas on the screen because individuals who do not look might be substantively different
488 than those that look.

489 We can fit a ZIB model using `brms()` and use the `{marginaleffects}` package to make
490 inferences about our parameters of interest. Before we run a zero-inflated beta model, we will
491 need to transform our data again and remove the one 1 value in our data—we can keep our 0s.
492 Similar to our beta regression model we fit in `brms`, we will use the `bf()` function to fit several
493 models. We fit our μ and ϕ parameters as well as our zero-inflated parameter (α ; here labeled as
494 `zi`). In `brms` we can use the `zero_inflated_beta` family (see Listing 15).

495 **Posterior Predictive Check**

496 The ZIB model does a bit better at capturing the structure of the data than the beta
497 regression model (see Figure 12). Specifically, the ZIB model more accurately captures the
498 increased density of values near the lower end of the scale (i.e., near zero), which the standard
499 beta model underestimates. The ZIB model's predictive distributions also align more closely with
500 the observed data across the entire range, particularly in the peak and tail regions. This improved
501 fit likely reflects the ZIB model's ability to explicitly model excess 0s (or near-zero values) via its
502 inflation component, allowing it to better account for features in the data that a standard beta
503 distribution cannot accommodate.

504 **Predicted Probabilities and Marginal Effects**

505 Table 8, under the zero-inflated beta regression column, provides a summary of the
506 posterior distribution for each parameter. As stated before, it is preferable to back-transform our
507 estimates to get probabilities. To get the predicted probabilities we can again use the

Listing 15 Fitting zib model with brm()

```
# keep 0 but remove 1

data_beta_0 <- fluency_data |>
  filter(Accuracy != 1)

# set up model formula for zero-inflated beta in brm

zib_model <- bf(
  Accuracy ~ Fluency, # The mean of the 0-1 values, or mu
  phi ~ Fluency, # The precision of the 0-1 values, or phi
  zi ~ Fluency, # The zero-or-one-inflated part, or alpha
  family = zero_inflated_beta()
)

# fit zib model with brm

fit_zi <- brm(
  formula = zib_model,
  data = data_beta_0,
  file = here::here("manuscript", "models", "bayes_zib_model0not1.rds")
)
```

508 avg_predictions() and avg_comparisons() functions from {marginaleffects} package
 509 (Arel-Bundock, 2024) to get predicted probabilities and the probability difference between the
 510 levels of each factor. We can model the parameters separately using the dpar argument setting to:
 511 μ , ϕ , α . Here we look at the risk difference for Fluency under each parameter. If one were
 512 interested in the average effect for the entire model, the dpar argument could be removed.

513 **Mu.** As shown in Table 4, there is little evidence for an effect of Fluency – the 95% Cr.I
 514 includes zero, suggesting substantial uncertainty about the direction and magnitude of the

515 effect—that is, though most of the posterior density supports positive effects, nil and weakly
 516 negative effects cannot be ruled out.

Table 4*Probability fluency difference (μ)*

| Contrast | Mean | 95% Cr.I | pd |
|--------------------|-------|-----------------|-------|
| Fluent - Disfluent | 0.043 | [-0.031, 0.118] | 0.875 |

517 **Dispersion.** As shown in Table 5, the posterior estimates suggest a credible effect of
 518 Fluency on dispersion (ϕ), with disfluent responses showing greater variability. The 95% Cr.I for
 519 the fluency contrast does not include zero, indicating a high probability in differences in precision.

Table 5*Probability fluency difference (ϕ)*

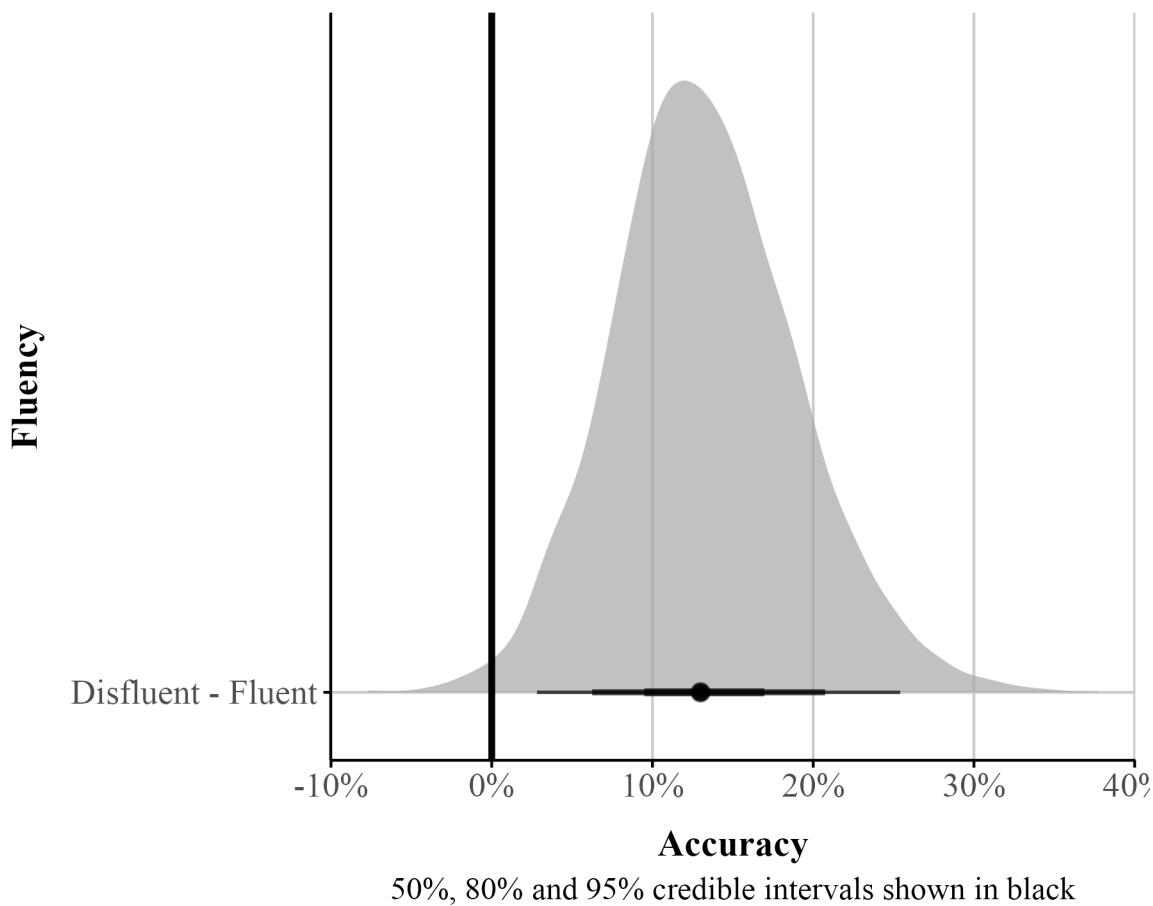
| Contrast | Mean | 95% Cr.I | pd |
|--------------------|------|----------------|-------|
| Fluent - Disfluent | 2.68 | [-0.894, 6.45] | 0.926 |

520 **Zero-Inflation**

521 We can use {marginaleffects} to estimate and plot the posterior difference between the
 522 fluent and disfluent conditions (see Figure 8). In Figure 8, the posterior distribution for this
 523 contrast lies mostly below zero, indicating that a fluent instructor is associated with a lower
 524 probability of zero responses. The estimated reduction is approximately 13%. The 95% credible
 525 interval does not include zero, which indicates that the data provide consistent evidence for a
 526 reduction in zero responses under fluent instruction.

Figure 8

Visualization of the predicted difference for zero-inflated part of model



527 **Zero-One-Inflated beta (ZOIB)**

528 The ZIB model works well if there are 0s in your data, but not 1s.⁸ In our previous
 529 examples we either got rid of both 0s and 1s (beta regression), or removed the 1s (ZIB).
 530 Sometimes it is theoretically useful to model both 0s and 1s as separate processes or to consider

⁸ In cases where your data include exact 1s but no 0s, you can fit a one-inflated beta regression model in {brms} by setting the `coi` parameter to 1. This tells the model that all point masses occur at 1, rather than being split between 0 and 1. In other words, `coi = 1` assumes that any inflation in the data is due entirely to values at 1. In our data, we have exactly one value equal to 1[⁶]. While probably not significant to alter our findings, we can model 1s with a special type of model called the zero-one-inflated beta (ZOIB) model (Liu & Kong, 2015) if we believe that both 0s and 1s are distinct outcomes.

531 these values as essentially similar parts of the continuous response, as we show later in the
 532 ordered beta regression model. For example, this is important in visual analog scale data where
 533 there might be a prevalence of responses at the bounds (Kong & Edwards, 2016), in JOL tasks
 534 (Wilford et al., 2020), or in a free-list task where individuals provide open responses to some
 535 question or topic which are then recoded to fall between 0-1 (Bendixen & Purzycki, 2023). Here
 536 0s and 1s are meaningful; 0 means item was not listed and 1 means the item was listed first.

537 Similar to the beta and zero-inflated models discussed above, we can fit a
 538 zero-and-one-inflated beta (ZOIB) model in {brms} using the `zero_one_inflated_beta` family.
 539 This formulation simultaneously estimates the mean μ and precision ϕ of the Beta component, as
 540 well as two inflation parameters: α , the probability that an observation is at either boundary (0 or
 541 1), and γ , the conditional probability that, given an observation falls on a boundary, it takes the
 542 value 1 rather than 0. In other words, α determines how often responses occur exactly at the
 543 endpoints, and γ determines the balance between zeros and ones among those endpoint values.
 544 This specification allows the model to capture both the continuous variation in the interior of the
 545 (0, 1) interval and the presence of exact boundary values.

546 To illustrate how α and γ shape the distribution, Figure 9 displays simulated data across a
 547 range of parameter combinations. As α increases, more responses occur at the endpoints. As γ
 548 increases, the proportion of those endpoint responses that are 1 increases relative to 0, producing
 549 increasingly pronounced spikes at 1 as γ approaches 1. Together, these parameters give the ZOIB
 550 model the flexibility to represent datasets with mixtures of continuous values and exact zeros and
 551 ones.

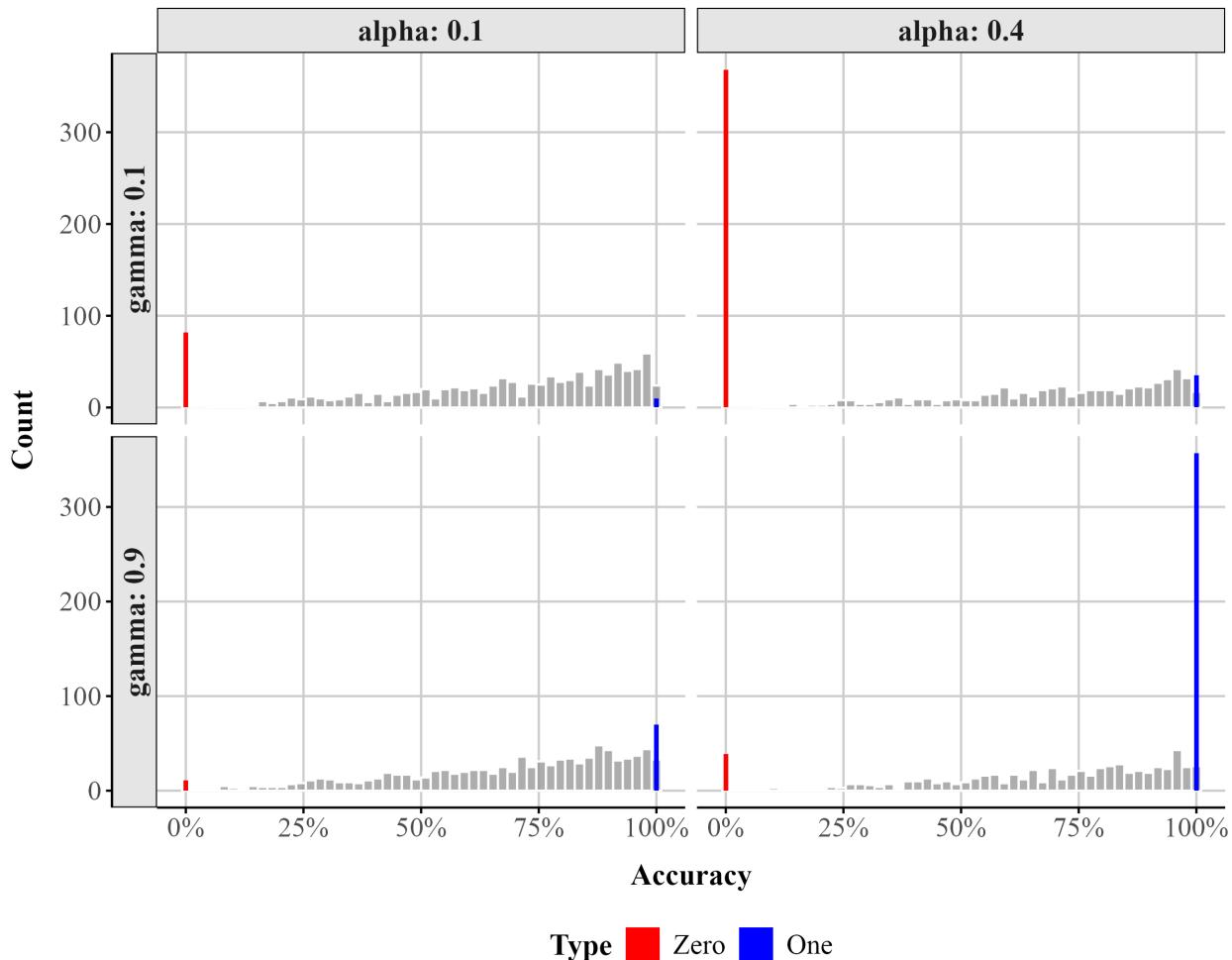
552 To fit a ZOIB model we use the `bf()` function. We model each parameter as a function of
 553 Fluency. We then pass the `zoib_model` to our `brm()` function (see Listing 16). The summary of
 554 the output is in Table 8 (under ZOIB).

555 **Model Parameters**

556 The output for the model is lengthy because we are estimating four distinct components,
 557 each with their own independent responses and sub-models. All the coefficients are on the logit

Figure 9

Simulated data from a ZOIB model illustrating the effects of the zero-one inflation parameter (α) and the conditional one-inflation parameter (γ).



558 scale, except ϕ , which is on the log scale. Thankfully drawing inferences for all these different
 559 parameters, plotting their distributions, and estimating their average marginal effects looks exactly
 560 the same—all the `brms` and `{marginaleffects}` functions we used work the same.

561 **Predictions and Marginal Effects**

562 With `{marginaleffects}` we can choose `marginalize` over all the sub-models, averaged
 563 across the 0s, continuous responses, and 1s in the data, or we can model the parameters separately
 564 using the `dpar` argument like we did above setting it to: μ , ϕ , α , γ (see below). Using

Listing 16 Fitting a ZOIB model with `brm()`.

```
# fit the zoib model

zoib_model <- bf(
  Accuracy ~ Fluency, # The mean of the 0-1 values, or mu
  phi ~ Fluency, # The precision of the 0-1 values, or phi
  zoi ~ Fluency, # The zero-or-one-inflated part, or alpha
  coi ~ Fluency, # The one-inflated part, conditional on the 1s, or gamma
  family = zero_one_inflated_beta()
)

fit_zoib <- brm(
  formula = zoib_model,
  data = fluency_data,
  file = here::here("manuscript", "models", "bayes_zoib_model")
)
```

565 avg_predictions() and not setting dpar we can get the predicted probabilities across all the
 566 sub-models. We can also plot the overall difference between fluency and disfluency for the whole
 567 model with plot_predictions().

568 In addition, we show below how one can extract the predicted probabilities and marginal
 569 effects for γ (and a similar process for any other model component, zoi, etc.):

Listing 17 Extracting predicted probabilities and marginal effects for conditional-one parameter

```
# get average predictions for coi param

coi_probs <- avg_predictions(fit_zoib, variables = "Fluency", dpar = "coi")
# get differene between the two conditions

coi_me <- avg_comparisons(fit_zoib, variables = c("Fluency"), dpar = "coi")
```

570 Ordered Beta Regression

571 Looking at the output from the ZOIB model (Table 8), we can see how running a model
572 like this can become fairly complex as it is fitting distinct sub-models for each component of the
573 scale. The ability to consider 0s and 1s as distinct processes from continuous values comes at a
574 price in terms of complexity and interpretability. A simplified version of the zero-one-inflated
575 beta (ZOIB) model, known as ordered beta regression (Kubinec, 2022; see also Makowski et al.,
576 2025 for a reparameterized version called the *beta-Gate* model), has been recently proposed. The
577 ordered beta regression model exploits the fact that, for most analyses, the continuous values
578 (between 0-1) and the discrete outcomes (e.g., 0 or 1) are *ordered*. For example, as a covariate x
579 increases or decreases, we should expect the bounded outcome y to increase or decrease
580 monotonically as well from 0 to (0, 1) to 1. The ZOIB model does not impose this restriction; a
581 covariate could increase and the response y could increase in its continuous values while
582 simultaneously decreasing at *both* end points.⁹ This complexity is not immediately obvious when
583 fitting the ZOIB, nor is it a potential relationship that many scholars want to consider when
584 examining how covariates influence a bounded scale.

585 To make the response ordered, the ordered beta regression model estimates a weighted
586 combination of a standard beta regression model for continuous responses and a logit model for
587 the discrete values of the response. By doing so, the amount of distinctiveness between the
588 continuous responses and the discrete end points is a function of the data (and any informative
589 priors) rather than strictly defined as fully distinct processes as in the ZOIB. For some datasets,
590 the continuous and discrete responses will be fairly distinct, and in others less so.

591 The weights that average together the two parts of the outcome (i.e., discrete and
592 continuous) are determined by cutpoints that are estimated in conjunction with the data in a
593 similar manner to what is known as an ordered logit model. An in-depth explanation of ordinal
594 regression is beyond the scope of this tutorial (Bürkner & Vuorre, 2019; but see Fullerton &
595 Anderson, 2021). At a basic level, ordinal regression models are useful for outcome variables that

⁹ For a more complete description of this issue, we refer the reader to Kubinec (2022).

596 are categorical in nature and have some inherent ordering (e.g., Likert scale items). To preserve
597 this ordering, ordinal models rely on the cumulative probability distribution. Within an ordinal
598 regression model it is assumed that there is a continuous but unobserved latent variable that
599 determines which of k ordinal responses will be selected. For example on a typical Likert scale
600 from ‘Strongly Disagree’ to ‘Strongly Agree’, you could assume that there is a continuous,
601 unobserved variable called ‘Agreement’.

602 While we cannot measure Agreement directly, the ordinal response gives us some
603 indication about where participants are on the continuous Agreement scale. $k - 1$ cutoffs are then
604 estimated to indicate the point on the continuous Agreement scale at which your Agreement level
605 is high enough to push you into the next ordinal category (say Agree to Strongly Agree).
606 Coefficients in the model estimate how much different predictors change the estimated *continuous*
607 scale (here, Agreement). Since there’s only one underlying process, there’s only one set of
608 coefficients to work with (proportional odds assumption).

609 In an ordered beta regression, three ordered categories are modeled: (1) exactly zero, (2)
610 somewhere between zero and one, and (3) exactly one. In an ordered beta regression, (1) and (2)
611 are modeled with cumulative logits, where one cutpoint is the the boundary between Exactly 0
612 and Between 0 and 1 and the other cutpoint is the boundary between *Between 0 and 1* and *Exactly*
613 *1*. The continuous values in the middle, 0 to 1 (3), are modeled as a vanilla beta regression with
614 parameters reflecting the mean response on the logit scale as we have described previously.
615 Ultimately, employing cutpoints allows for a smooth transition between the bounds and the
616 continuous values, permitting both to be considered together rather than modeled separately as the
617 ZOIB requires.

618 The ordered beta regression model has shown to be more efficient and less biased than
619 some of the methods discussed (Kubinec, 2022) herein and has seen increasing use across the
620 biomedical and social sciences (Martin et al., 2024; Nouvian et al., 2023; Shrestha et al., 2024;
621 Smith et al., 2024; Wilkes et al., 2024) because it produces only a single set of coefficient

622 estimates in a similar manner to a standard beta regression or OLS.¹⁰

623 ***Fitting an Ordered Beta Regression***

624 To fit an ordered beta regression in a Bayesian context we use the `{ordbetareg}` (Kubinec,
 625 2023) package. `{ordbetareg}` is a front-end to the `brms` package that we described earlier; in
 626 addition to the functions available in the package, most `brms` functions and plots, including the
 627 diverse array of regression modeling options, will work with `{ordbetareg}` models. (We note that
 628 the `ordbeta` model is also available as a maximum-likelihood variant in the R package
 629 `{glmmTMB}`.) We first load the `{ordbetareg}` package (see Listing 18).

Listing 18 Load `{ordbetareg}`

```
library(ordbetareg)
```

630 The `{ordbetareg}` package uses `brms` on the front-end so all the arguments we used
 631 previously apply here. Instead of the `brm()` function we use `ordbetareg()`. To fit a model where
 632 dispersion does not vary as a function of fluency we can use the below code (see Listing 19).

Listing 19 Fitting ordered beta model with `ordbetareg()`

```
ord_fit_brms <- ordbetareg(  

  Accuracy ~ Fluency,  

  data = fluency_data,  

  file = here::here("manuscript", "models", "bayes_ordbeta_model")  

)
```

633 However, if we want dispersion to vary as a function of fluency we can easily do that (see
 634 Listing 20). Note the addition of the `phi_reg` argument in `m.phi`. This argument allows us to

¹⁰ Please note that there are other models available that can model this continuous process like the beta-gate model (Makowski et al., 2025) and the censored extended beta regression model (Kosmidis & Zeileis, 2025).

- 635 include a model that explicitly models the dispersion parameter. Because we are modeling ϕ as a
 636 function of fluency, we set the the argument to both.

Listing 20 Fitting ordered beta model with dispersion using `ordbetareg()`

```
ord_beta_phi <- bf(Accuracy ~ Fluency, phi ~ Fluency)

m.phi <- ordbetareg(
  ord_beta_phi,
  data = fluency_data,
  phi_reg = 'both',
  file = here::here("manuscript", "models", "bayes_ordbeta_phi_model")
)
```

- 637 **Marginal Effects.** Table 8 presents the posterior summary. We can use {marginaleffects}
 638 to calculate differences on the response scale that average over (or marginalize over) all our
 639 parameters.

Table 6

Marginal effect of fluency ordered beta model

| Contrast | Mean | 95% Cr.I | pd |
|--------------------|-------|-----------------|------|
| Fluent - Disfluent | 0.061 | [-0.016, 0.139] | 0.94 |

- 640 In Table 6 the credible interval is close enough to zero relative to its uncertainty that we
 641 can conclude there likely aren't differences between the conditions after taking dispersion and the
 642 0s and 1s in our data into account.

- 643 **Cutpoints.** The model cutpoints are not reported by default in the summary output, but
 644 we can access them with the R package `posterior` (Bürkner et al., 2025) and the functions
 645 `as_draws` and `summary_draws`.

Table 7

Cutzero and cutone parameter summary

| Parameter | Mean | 95% Cr.I |
|-----------|-------|----------------|
| cutzero | -2.97 | [-3.57, -2.44] |
| cutone | 1.85 | [1.64, 2.07] |

In Table 7, cutzero is the first cutpoint (the difference between 0 and continuous values)

and cutone is the second cutpoint (the difference between the continuous values and 1). These cutpoints are on the logit scale and as such the numbers do not have a simple substantive meaning.

In general, as the cutpoints increase in absolute value (away from zero), then the discrete/boundary observations are more distinct from the continuous values. This will happen if there is a clear gap or bunching in the outcome around the bounds. This type of empirical feature of the distribution may be useful to scholars if they want to study differences in how people perceive the ends of the scale versus the middle. It is possible, though beyond the scope of this article, to model the location of the cutpoints with hierarchical (non-linear) covariates in brms. In the most recent version of ordbeta, it is possible to test the influence of different factors on these boundaries.

Model Fit

The best way to visualize model fit is to plot the full predictive distribution relative to the original outcome. Because ordered beta regression is a mixed discrete/continuous model, a separate plotting function, pp_check_ordbetareg, is included in the {ordbetareg} package that accurately handles the unique features of this distribution. The default plot in brms will collapse these two features of the outcome together, which will make the fit look worse than it actually is.

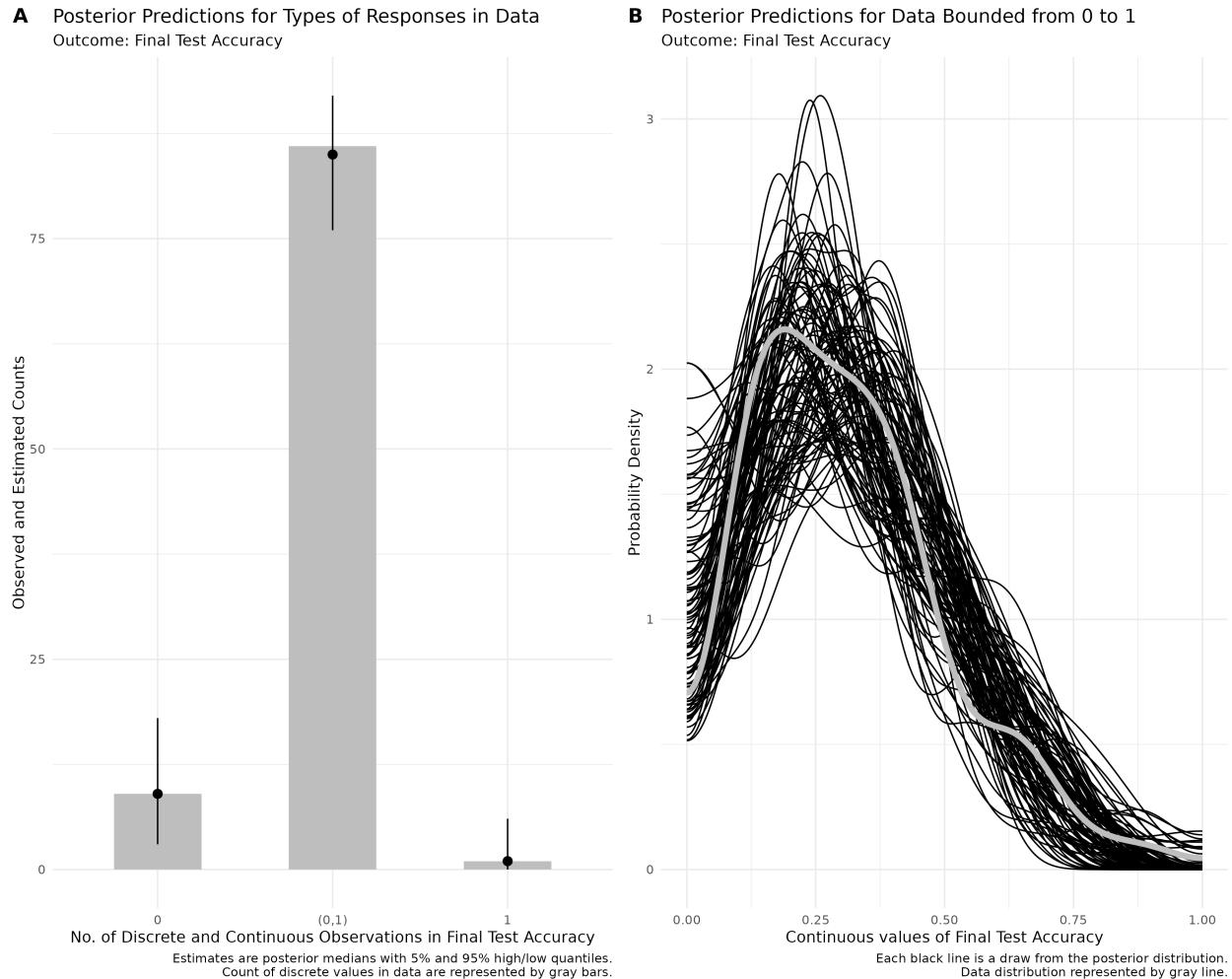
The {ordbetareg} function returns a list with two plots, discrete and continuous, which can either be printed and plotted or further modified as {ggplot2} objects (see Figure 10).

The discrete plot, which is a bar graph, shows that the posterior distribution accurately

captures the number of different types of responses (discrete or continuous) in the data. For the

Figure 10

Posterior predictive check for ordered beta regression model. A. Discrete posterior check. B. Continuous posterior check.



666 continuous plot shown as a density plot with one line per posterior draw, the model does a very
 667 good job at capturing the distribution.

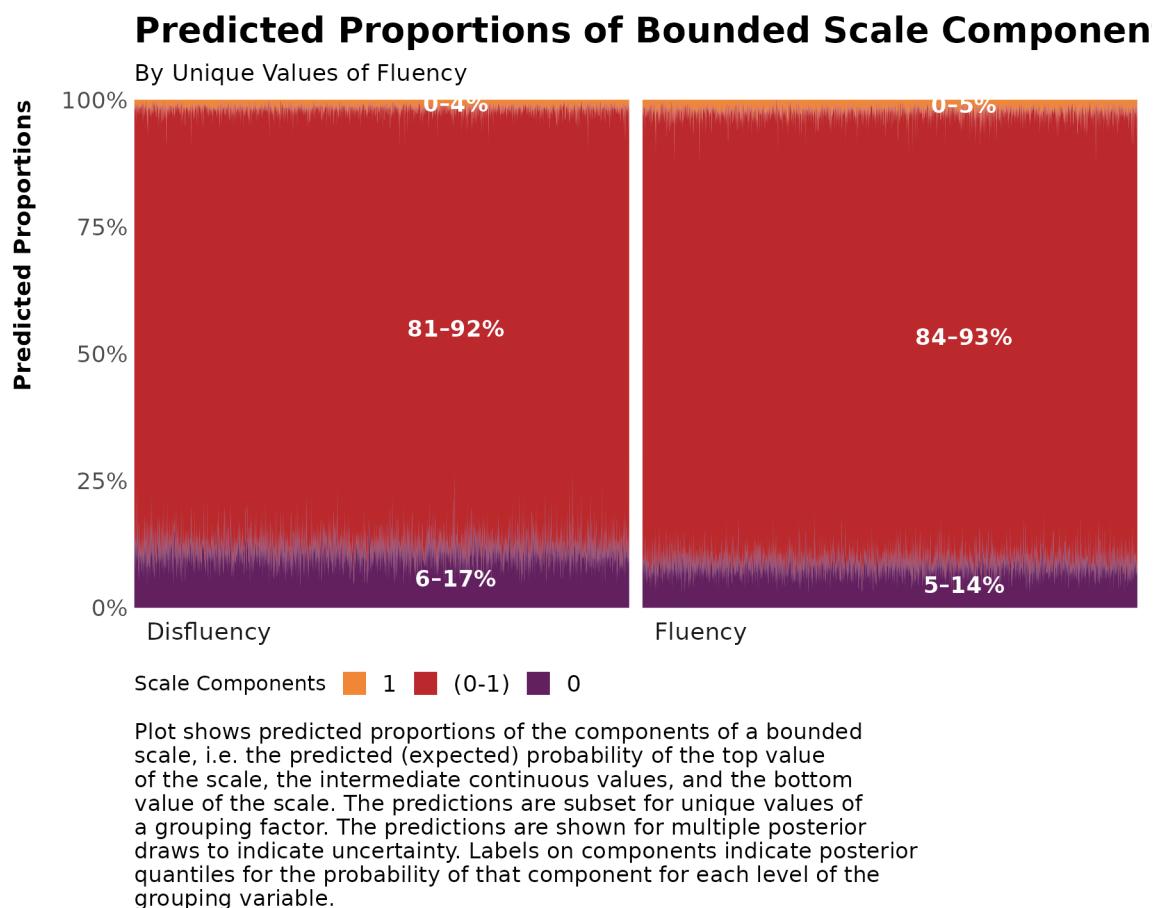
668 Overall, it is clear from the posterior distribution plot that the ordered beta model fits the
 669 data well. To fully understand model fit, both of these plots need to be inspected as they are
 670 conceptually distinct.

671 ***Model Visualization***

672 {ordbetareg} provides a useful visualization function called `plot_heiss()` (Ye & Heiss,
 673 2023) that can represent dispersion in the entire outcome as a function of discrete covariates. This
 674 function produces a plot of predicted proportions across the range of our Fluency factor. In
 675 Figure 11 we get predicted proportions for Fluency across the bounded scale. Looking at the
 676 figure we can see there is much overlap between instructors in the middle portion (μ) . However,
 677 we do see some small differences at the zero bounds.

Figure 11

Heiss plot of predicted probabilities across the scale (0-100)

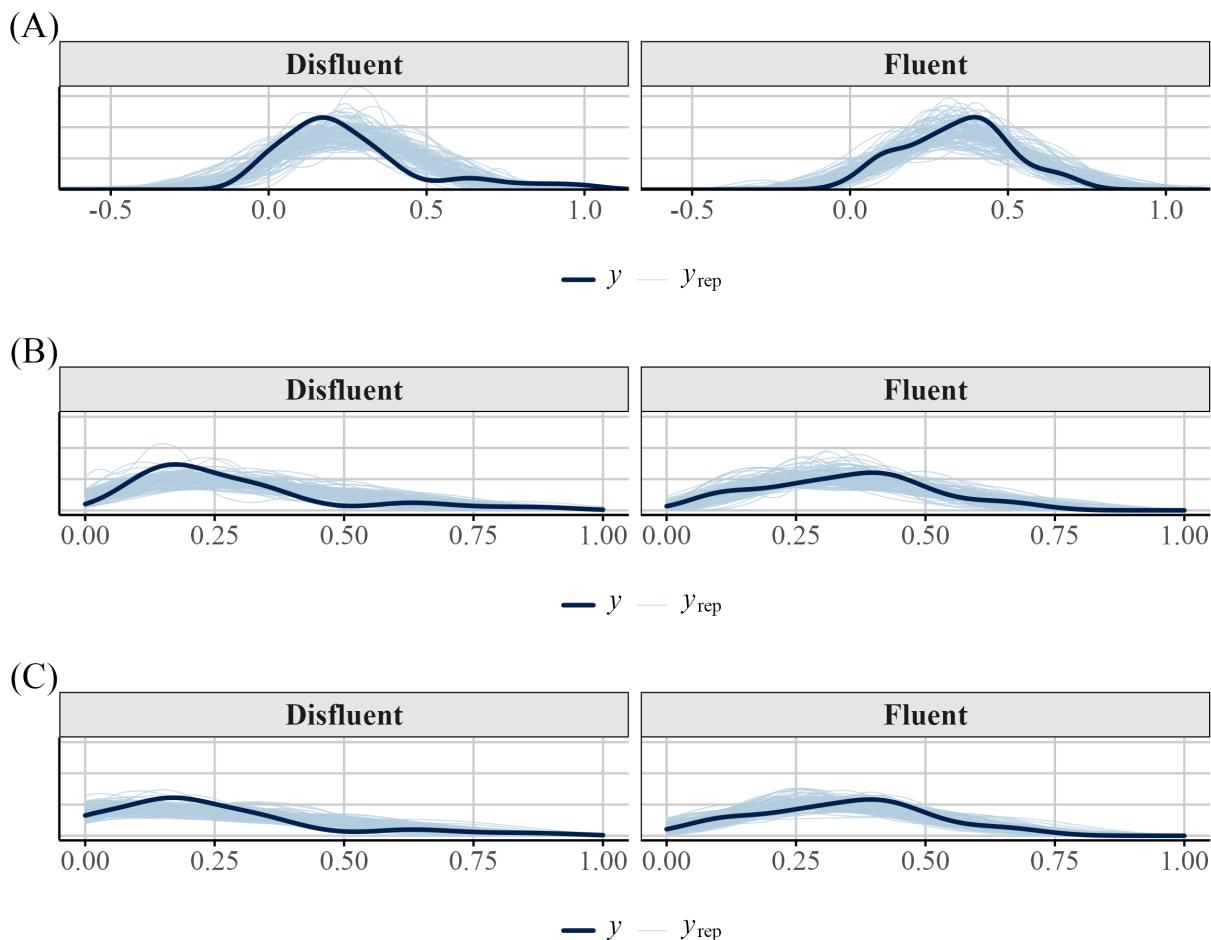


678 **Ordered Beta Scale**

679 In the `{ordbetareg}` function there is a `true_bound` argument. In cases where your data is
 680 not bounded between 0-1, this argument can be used to specify the bounds of the argument to fit
 681 the ordered beta regression. For example, the response data might be bounded between 1 and 7. If
 682 so, `{ordbetareg}` can model it within the [0,1] interval and `{ordbetareg}` will convert the model
 683 predictions back to the true bounds after estimation.

Figure 12

The plots show 100 posterior predicted distributions with the label y_{rep} (light blue), the distribution of accuracy as function of fluency in dark blue for regular regression (A), beta regression (B), ZIB (C) and ZOIB (D).



684

Discussion

685 The use of beta regression in psychology, and the social sciences in general, is rare. With
686 this tutorial, we hope to turn the tides. Beta regression models are an attractive alternative to
687 models that impose unrealistic assumptions like normality, linearity, homoscedasticity, and
688 unbounded data. Beyond these models, there are a diverse array of different models that can be
689 used depending on your outcome of interest.

690 Throughout this tutorial our main aim was to help guide researchers in running analyses
691 with proportional or percentage outcomes using beta regression and some of its alternatives. In
692 the current example, we used real data from Wilford et al. (2020) and discussed how to fit these
693 models in R, interpret model parameters, extract predicted probabilities and marginal effects, and
694 visualize the results using the `{marginaleffects}` package.

695 Comparing our analysis with that of Wilford et al. (2020), we demonstrated that using a
696 traditional approach (e.g., *t*-test) to analyze mean accuracy data can lead to biased inferences.
697 Although we successfully reproduced one of their key findings, our use of beta regression and its
698 extensions revealed important nuances in the results. With a traditional beta regression
699 model—which accounts for both the mean and the precision (dispersion)—we observed similar
700 effects of instructor fluency on performance. However, the standard beta model does not
701 accommodate boundary values (i.e., 0s and 1s).

702 When we applied a ZIB model, which explicitly accounts for structural 0s, we found no
703 effect of fluency on the mean (μ) part of the model. Instead, the effect of fluency emerged in the
704 structural zero (inflated zero; α) component. This pattern was consistent when using a
705 zero-one-inflated beta (ZOIB) model. Furthermore, we fit an ordered beta regression model
706 (Kubinec, 2022), which appropriately models the full range of values, including 0s and 1s. Here,
707 we did not observe a reliable effect of fluency on the mean once we accounted for dispersion.

708 These analyses emphasize the importance of fitting a model that aligns with the nature of
709 the data. The simplest and recommended approach when dealing with data that contains 0s and/or
710 1s is to fit an ordered beta model, assuming the process is truly continuous. However, if you

711 believe the process is distinct in nature, a ZIB or ZOIB model might be a better choice.

712 Ultimately, this decision should be guided by theory.

713 For instance, if we believe fluency influences the boundaries (0 and 1), we might want to
714 model this process separately using a ZIB or ZOIB. With the current dataset, fluency might affect
715 specific aspects of performance (such as the likelihood of complete failure) rather than general
716 performance levels. This effect could be due to participant disengagement during the disfluent
717 lecture. If students fail to pay attention because of features of disfluency, they may miss relevant
718 information, leading to a floor effect at the test. Following from this, disfluency would be
719 expected to influence the boundary (0) and not the continuous part of the model. If this is the
720 case, we would want to model this appropriately. However, if we believe fluency effects general
721 performance levels (the continuous part), a model that takes in to account the entire process
722 accounting for the 0s and 1s might be appropriate.

723 In the discussion section of Wilford et al. (2020), they were unable to offer a tenable
724 explanation for performance differences based on instructor fluency. A model that accounts for the
725 excess 0s in the dataset provides one testable explanation: watching a disfluent lecture may lead to
726 lapses in attention, resulting in poorer performance in that group. These lapses, in turn, contribute
727 to the observed differences in the fluent condition. This modeling approach opens a promising
728 avenue for future research—one that would have remained inaccessible otherwise.

729 Not everyone will be eager to implement the techniques discussed herein. In such cases,
730 the key question becomes: What is the least problematic approach to handling proportional data?
731 One reasonable option is to fit multiple models tailored to the specific characteristics of your data.
732 For example, if your data contain 0s, you might fit two models: a traditional linear model
733 excluding the 0s, and a logistic model to account for the zero versus non-zero distinction. If your
734 data contain both 0s and 1s, you could fit separate models for the 0s and 1s in addition to the OLS
735 model. There are many defensible strategies to choose from depending on the context. However,
736 we do not recommend transforming the values of your data (e.g., 0s to .01 and 1s to .99) or
737 ignoring the properties of your data simply to fit traditional statistical models.

738 In this tutorial, we demonstrated how to analyze these models from a Bayesian perspective.

739 While we recognize that not everyone identifies as a Bayesian, implementing these models using a
740 Bayesian framework is relatively straightforward—it requires only a single package, lowering the
741 barrier to entry. For those who prefer frequentist analyses, several R packages are available. For
742 example, the `{betareg}` package (Cribari-Neto & Zeileis, 2010) `{glmmTMB}` (Brooks et al.,
743 2017) and `{gamlss}` (2005) are nice options. To this end, I have included supplemental materials
744 demonstrating how to use frequentist packages to analyze the data presented herein.

745 **Conclusion**

746 Overall, this tutorial emphasizes the importance of modeling the data you have. Although

747 the example provided is relatively simple (a one-factor model with two levels), we hope it
748 demonstrates that even with a basic dataset, there is much nuance in interpretation and inference.
749 Properly modeling your data can lead to deeper insights, far beyond what traditional measures
750 might offer. With the tools introduced in this tutorial, researchers now have the means to analyze
751 their data effectively, uncover patterns, make accurate predictions, and support their findings with
752 robust statistical evidence. By applying these modeling techniques, researchers can improve the
753 validity and reliability of their studies, ultimately leading to more informed decisions and
754 advancements in their respective fields.

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Table 8*Bayesian regression summaries for each model*

| Parameter | Stat | Bayesian | Beta | ZIB | ZOIB | Ordered |
|-----------------|------|----------------|------------------|------------------|------------------|------------------|
| | | LM | Regression | | | Beta |
| b_Intercept | Mean | 0.256 | -0.831 | -0.828 | -0.826 | -0.865 |
| | Cr.I | [0.201, 0.313] | [-1.102, -0.561] | [-1.1, -0.564] | [-1.093, -0.556] | [-1.124, -0.606] |
| | pd | 1.000*** | 1.000*** | 1.000*** | 1.000*** | 1.000*** |
| b_Fluency | Mean | 0.085 | 0.201 | 0.198 | 0.198 | 0.261 |
| | Cr.I | [0, 0.167] | [-0.156, 0.552] | [-0.14, 0.543] | [-0.151, 0.54] | [-0.069, 0.597] |
| | pd | 0.976* | 0.871 | 0.875 | 0.870 | 0.940 |
| sigma | Mean | 0.208 | - | - | - | - |
| | Cr.I | [0.182, 0.24] | - | - | - | - |
| | pd | 1.000*** | - | - | - | - |
| b_phi_Intercept | Mean | - | 1.599 | 1.603 | 1.600 | 1.619 |
| | Cr.I | - | [1.172, 1.986] | [1.183, 1.984] | [1.176, 1.985] | [1.204, 1.996] |
| | pd | - | 1.000*** | 1.000*** | 1.000*** | 1.000*** |
| b_phi_Fluency | Mean | - | 0.429 | 0.424 | 0.434 | 0.396 |
| | Cr.I | - | [-0.157, 0.996] | [-0.155, 0.993] | [-0.131, 1.021] | [-0.202, 0.966] |
| | pd | - | 0.928 | 0.926 | 0.929 | 0.907 |
| b_zi_Intercept | Mean | - | - | -1.670 | - | - |
| | Cr.I | - | - | [-2.476, -0.953] | - | - |